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Perfect score:
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                                                                                                                                                                                                                                                                                                                                          Score
length: 0
length: 2000000000
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24
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1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*

2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*

3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*

4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*

5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*

6: /cgn2_6/ptodata/1/ina/backfiles1.seq:*
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Maximum Match 100%
Listing first 45 summaries
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2 US-08-439-814-3

2 US-08-439-814-1

2 US-08-439-814-1

3 US-09-949-016-70889

4 US-09-949-016-1186

6 US-09-949-016-1186

6 US-09-621-976-17501

6 US-09-621-976-17501

7 US-09-621-976-17501

8 US-09-621-976-17502

8 US-09-621-976-17502

9 US-09-621-976-13989

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9 US-09-949-016-16417

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Compugen Ltd
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Sequence 3, Appli
Sequence 2, Appli
Sequence 1, Appli
Sequence 70889, A
Sequence 70890, A
Sequence 49, Appl
Sequence 11186, A
           Sequence
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                                                                                 9575, Ap
16117, A
12422, A
14718, A
29459, A
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17502, A
13989, A
11249, A
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13840, A
                                    195084
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ALIGNMENTS

US-08-439-814-4/c RESULT 1 COUNTRY 2005-5701 ZIE: 2005-5701 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS OPERATING SYSTEM: PC-DOS/MS-DOS OPERATING SYSTEM: PC-DOS/MS-DOS OPERATING SYSTEM: PC-DOS/MS-DOS OURRENT APPLICATION NATA: APPLICATION NUMBER: US/08/439,814 PRIOR APPLICATION DATA: APPLICATION DATA: APPLICATION DATA: APPLICATION NUMBER: US P 4238778.7 FILING DATE: 12-NOV-1992 PRIOR APPLICATION DATA: APPLICATION NUMBER: DE PCT/DE93/01086 FILING DATE: 10-NOV-1992 PRIOR APPLICATION NUMBER: DE PCT/DE93/01086 FILING DATE: 10-NOV-1993 ATTORNEY/AGENT INFORMATION: NAME: KLESNER, Sharon N. REFERENCE/DOCKET NUMBER: D1614-5015 REFERENCE/DOCKET NUMBER: D1614-5015 TELECOMMUNICATION INFORMATION: TELEPHONE: 202/638-4810 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: TRM PC COMPATIBLE COMPUTER: TRM PC COMPATIBLE TELECOMMUNICATION TRM PC COMPATIBLE COMPUTER: TRM PC COMPATIBLE Sequence 4, Application US/08439814 Patent No. 5968735 GENERAL INFORMATION: APPLICANT: STEIN, Ulrike APPLICANT: WALTHER, Wolfgang TITLE OF INVENTION: VECTOR FOR THE EXPRESSION TITLE OF INVENTION: THERAPY-RELEVANT GENES COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PATENTIN Release #1.0, VG PRIOR APPLICATION DATA: APPLICATION UNMBER: DE P4238778.7 FILING DATE: 12-NOV-1992 PRIOR APPLICATION DATA: NUMBER OF SEQUENCES: 7 CORRESPONDENCE ADDRESS: ADDRESSEE: NIKAIDO, MARMELSTEIN, MURRAY & ORAM STREET: 655 Fifteenth Street, N. W., Suite 330 STREET: Street Lobby CITY: Washington APPLICATION NUMBER: USA WO PCT/DE93/01086 Version #1.30B ဝှု (EPO)

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                                                                                                                                                                                                                                                                                 CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE P 4238778.7
FILLING DATE: 12-NOV-1992
PRIOR APPLICATION NUMBER: DE PCT/DE93/01086
FILLING DATE: 10-NOV-1993
ATTORNEY/AGENT INFORMATION:
NAME: KLESNER, Sharon N.
REFERENCE/DOCKET NUMBER: 91614-5015
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202/638-5000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 3, Application US/08439814
Patent No. 5968735
GENERAL INFORMATION:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PRILING DATE: 10-NOV-1993
NFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
                                                                                              COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30B (EPO)
PRIOR APPLICATION DATA;
APPLICATION NUMBER: DE P4238778.7
FILING DATE: 12-NOV-1992
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MEDIUM TYPE: Floppy disk
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APPLICATION NUMBER: US
FILING DATE: 12-MAY-19
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MEDIUM TYPE: Floppy disk
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APPLICANT: WALTHER, Wolfgang
TITLE OF INVENTION: VECTOR FOR THE EXPRESSION OF
TITLE OF INVENTION: THERAPY-RELEVANT GENES
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LENGTH: 568 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 655 Fifteenth Street, N. W., Suite 330 G STREET: Street Lobby CITY: Washington STATE: DC
                                                                                                                                                                                                                                                              TELEPHONE: 202/638-500
TELEFAX: 202/638-4810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING PATENTIN Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      / Match 100.0%; Score 24; DB Local Similarity 100.0%; Pred. No. 0.02 nes 24; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
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                                                             WO PCT/DE93/01086
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/439,814
FILING DATE: 12-MXY-1995
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION UNMBER: DE P4238778.7
FILING DATE: 12-MOV-1992
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION UNMBER: DE P67/DE93/01086
FILING DATE: 10-MOV-1993
ATTORNEY/AGENT INFORMATION:
NAME: KIRENEE SPACE N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 2, Application US/08439814 Patent No. 5968735 GENERAL INFORMATION:
                                                                                             APPLICATION NUMBER: WO FILING DATE: 10-NOV-19 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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Best Local Similarity 100.0%;
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve.
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE P4238778.7
FILING DATE: 12-NOV-1992
PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                 NAME: KLESNER, Sharon N.
REGISTRATION NUMBER: 36,335
REFERENCE/DOCKET NUMBER: P161
TELECOMPUNICATION INFORMATION:
TELEPHONE: 202/638-5000
TELEFAX: 202/638-4810
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                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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STREET: Street Lobby
CITY: Washington
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: WALTHER, WOLIGANG
TITLE OF INVENTION: VECTOR FOR THE EXPRESSION OF
TITLE OF INVENTION: THERAPY-RELEVANT GENES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                    STRANDEDNESS: single
    COPOLOGY:
                                                                             LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
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                      linear
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EQ ID NO: 2:
                                                                                                                                                                                                                                                                                                             Floppy disk
DNA (genomic)
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100.0%; Pred. No. 0
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                                                       US-08-439-814-1
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Patent No. 5968735
GENERAL INFORMATION:
APPLICANT: STEIN, Ulrike
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
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Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/439,814
FILING DATE: 12-MAY-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: DE P 4238778.7
FILING DATE: 12-NOV-1992
PRIOR APPLICATION NUMBER: DE PCT/DE93/01086
FILING DATE: 10-NOV-1993
ATTORNEY/AGENT INFORMATION:
NAME: KLESNER, Sharon N.
REGISTRATION NUMBER: 36,335
REFERENCE/DOCKET NUMBER: 91614-5015
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202/638-5000
                                                                                                                                                        INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: STEIN, ULTIKE
APPLICANT: WALTHER, WOLFGANG
TITLE OF INVENTION: VECTOR FOR THE EXPRESSION OF
TITLE OF INVENTION: THERAPY-RELEVANT GENES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS;
CADDRESSEE: NIKAIDO, MARMELSTEIN,
STREET: 655 Fifteenth Street, N.
STREET: Street Lobby
                                                                                                                                                                                                               FILING DATE: 12-NOV-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PO
                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.30B PRIOR APPLICATION DATA: APPLICATION NUMBER: DE P4238778.7
                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: 202/638-500
TELEFAX: 202/638-4810
COMPUTER READABLE FORM:
                                                                        MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
                                                                                         TOPOLOGY:
                                                                                                        STRANDEDNESS: single
                                                                                                                                          LENGTH:
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                                                                                                                           nucleic acid
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                                                                                       linear
                                                                      DNA (genomic)
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                                                                                                                                                                                                                      WO PCT/DE93/01086
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Pred. No. 0.032;
 Score 24; DB 2;
Pred. No. 0.033;
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W., Suite 330 G
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                   Length 2090;
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; ORGANISM: Human
US-09-949-016-70889
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
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US-09-949-016-70890/c
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
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Patent No. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 70889
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 70890
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Best Local Similarity
                                                                     Best Local Similarity Matches 19; Conserv
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                                                                                                        Query Match
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: CL001307
                                                                                                                                                                  TYPE: DNA
ORGANISM: Human
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 TTATGTATACATGTCTTTTAAA 74
                                  TTATGTAGACACGTCTTTCAAA 23
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Pred. No. 61;
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Pred. No. 6
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Sequence 14186, Application US/09949016
PATENT NO. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
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Best Local Similarity 86.4%;
Matches 19; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Lander
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Granahan, Patricia
REGISTRATION NUMBER: 32,227
REFERENCE/DOCKET NUMBER: WH
TELECOMMUNICATION INFORMATION:
TELEPHONE: 781-861-6240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/09/054,272
FILING DATE: 01-APR-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM: Windows 95
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskett
COMPUTER: IBM Compat
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TITLE OF INVENTION: IN VASCULAR PATHOLOGY GENES
NUMBER OF SEQUENCES: 59
CORRESPONDENCE ADDRESS:
ADDRESSEE: HAMILTON, BROOK, SMITH & REYNOLDS, P.C.
STREET: Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 13204 E
TYPE: nucleic a
STRANDEDNESS: 1
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APPLICANT: Irela
APPLICANT: Rozen
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ucleic acid
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Daley, George Q.
Cargill, Michele
Ireland, James S.
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SYSTEM: Windows 95
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; LENGTH: 125536
; TYPE: DNA
; ORGANISM: Human
US-09-949-016-14186
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US-09-621-976-17500/c
                                                                                                                                                                                                                                                                                                                      문
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; OTHER INFORMATION: n = A, T, C or G
US-09-949-016-13840
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; Sequence 13840, Application US/09949016
; Patent No. 6812339
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                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                    Patent No. 6639063
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LENGTH: 276687
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APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Jobert, S.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: ESTS and Encoded Human Proteins.
FILE REFERENCE: GENSET.054PR2
CURRENT APPLICATION NUMBER: US/09/621,976
CURRENT APPLING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 19335
SOFTWARE: Patent.pm
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Matches 19; Conserv
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 14186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTMARE: FastSEQ for Windows Version 4.0
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APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

FILE REFERENCE: CL001307

CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR FILING DATE: 2000-10-20

PRIOR FILING DATE: 2000-10-20
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PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Human
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Pred. No. 1.
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Pred. No. 1.
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APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: ESTS and Encoded Human Proteins.
FILE REFERENCE: GENSET.054PR2
CURRENT APPLICATION NUMBER: US/09/621,976
CURRENT FILING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 19335
SOFTWARE: Patent.pm
SEQ ID NO 17501
LENGTH: 103
TYPE: DNA
ORGANISM: Homo sapiens
US-09-621-976-17501
                                                                                                                                                            APPLICANT: Dumas Wilne Edwards, J.B.
APPLICANT: Jobert, S.
APPLICANT: Glordano, J.Y.
APPLICANT: Glordano, J.Y.
FILE OF INVENTION: ESTS and Encoded Human Proteins.
FILE REFERENCE: GENSET.054PR2
CURRENT APPLICATION NUMBER: US/09/621,976
CURRENT FILING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 19335
SOFTWARE: Patent.pm
SEQ ID NO 17503
LENGTH: 126
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US-09-621-976-17501/c
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-621-976-17500
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                                                                                                                                                                                                                                                                                                                                                                                   US-09-621-976-17503/c
                                                                                                                     ; ORGANISM: Homo sapiens
US-09-621-976-17503
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                                                                                                                                                                                                                                                                                                                                   Sequence 17503, Application US/09621976
Patent No. 6639063
GENERAL INFORMATION:
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Best Local Similarity
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Matches 17; Conserv
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                                                                          Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Dumas Milne Edwards, J.B.
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                   1 CTTATGTAGACACGTCT 17
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17; Conserv
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   CTTATGTAGACACGTCT 58
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                                                                          70.8%; Score 17; DB 4; Length 126; 100.0%; Pred. No. 59;
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Sequence 11249, Application US/09621976

Patent No. 6639063

GENERAL INFORMATION:
APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Giordano, J.Y.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: ESTS and Encoded Human Proteins.
FILE REFERENCE: GENSET.054PR2
CURRENT APPLICATION NUMBER: US/09/621,976
CURRENT FILING DATE: 2000-07-21

NUMBER OF SEQ ID NOS: 19335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-621-976-17502
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US-09-621-976-11249/c
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US-09-621-976-13989/c
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PATENT NO. 6639063
GENERAL INFORMATION:
APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Jobert, S.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: ESTS and Encoded Human Proteins.
FILE REFERENCE: GENSET.054PR2
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SEQ ID NO 17502
LENGTH: 137
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CURRENT FILING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 19335
SOFTWARE: PAtent.pm
SEQ ID NO 13989
LENGTH: 150
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Patent No. 6639063
GENERAL INFORMATION:
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Best Local 9
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CURRENT FILING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 19335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Jobert, S.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: ESTs and Encoded Human Proteins.
FILE REFERENCE: GENSET.054PR2
SOFTWARE: Patent.pm
SEQ ID NO 11249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    70.8%; Score 17;
Local Similarity 100.0%; Pred. No.
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Conservative (
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Minimum DB seq length: 0
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Maximum Match 100%
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Published Applications NA:*

1: /cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*

2: /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*

3: /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*

4: /cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*

5: /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*

6: /cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*

7: /cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*

9: /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*

9: /cgn2_6/ptodata/1/pubpna/US09E_PUBCOMB.seq:*

10: /cgn2_6/ptodata/1/pubpna/US09E_PUBCOMB.seq:*

11: /cgn2_6/ptodata/1/pubpna/US09C_PUBCOMB.seq:*

13: /cgn2_6/ptodata/1/pubpna/US09C_PUBCOMB.seq:*

14: /cgn2_6/ptodata/1/pubpna/US10E_PUBCOMB.seq:*

15: /cgn2_6/ptodata/1/pubpna/US10E_PUBCOMB.seq:*

16: /cgn2_6/ptodata/1/pubpna/US10E_PUBCOMB.seq:*

17: /cgn2_6/ptodata/1/pubpna/US10E_PUBCOMB.seq:*

18: /cgn2_6/ptodata/1/pubpna/US10E_PUBCOMB.seq:*

19: /cgn2_6/ptodata/1/pubpna/US10E_PUBCOMB.seq:*

19: /cgn2_6/ptodata/1/pubpna/US10E_PUBCOMB.seq:*

19: /cgn2_6/ptodata/1/pubpna/US10E_PUBCOMB.seq:*

20: /cgn2_6/ptodata/1/pubpna/US10E_PUBCOMB.seq:*

21: /cgn2_6/ptodata/1/pubpna/US10E_PUBCOMB.seq:*
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Copyright (c) 1993 - 2005 Compugen Ltd.
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539.601 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	. Score	공성	Length DB	BB	ID	Description
o :	1 24	100.0	633	18	US-10-482-433A-3	Sequence 3, Appli
ი	2 24	100.0	2932	18	US-10-473-126-37	Sequence 37, Appl
ი	3 19.8	82.5	633	18	US-10-482-433A-6	Sequence 6, Appli
ი	4 19.8	82.5	2932	17	US-10-451-646-24	Sequence 24, Appl
a	5 19:8	82.5	2932	18	US-10-473-126-167	Sequence 167, App
	6 18.2	75.8	491	18	US-10-437-963-86300	Sequence 86300, A
ი	7 18.2	75.8	494	10	US-09-770-961-577	Sequence 577, App
	8 18.2	75.8	520	18	US-10-357-930-19156	Sequence 19156, A
ი	9 18.2	75.8	597	18	US-10-357-930-54995	Sequence 54995, A
_	0 18.2	75.8	642	18	US-10-357-930-48960	Sequence 48960, A
u	1 18.2	75.8	1504	18	US-10-437-963-86301	Sequence 86301, A

RESULT 2 US-10-473-126-37/c

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US-09-864-761-6452 US-10-637-855-398 US-10-027-632-214760 US-10-027-632-214761	-10-424-599-3577 -10-282-122A-183 -09-764-872-700 -10-719-993-6811 -10-097-111-10	10-027-632- 10-027-632- 10-027-632- 10-437-963- 10-097-111-	US-10-027-632-8632 US-10-027-632-9405 US-10-027-632-99405 US-10-027-632-99405 US-10-027-632-99405 US-10-027-632-99405 US-10-719-993-6890 US-09-814-353-5727 US-09-814-353-5727 US-09-814-353-5727 US-10-82-1722A-38309 US-10-741-601-1999 US-10-741-601-1999 US-10-741-601-1999 US-10-98-2638-45459 US-10-097-111-147 US-10-097-111-147 US-10-097-632-274932	US-10-473-126-168 US-10-473-126-313
398 214 214	35770, 18304, 700, Ap 6811, A		Sequence 8632, Ap Sequence 8632, Ap Sequence 99405, A Sequence 99405, A Sequence 99405, A Sequence 99406, A Sequence 5727, Ap Sequence 12011, A Sequence 12011, A Sequence 50147, A Sequence 5959, A Sequence 45459, A Sequence 147, App Sequence 147, App Sequence 147, App Sequence 147, App Sequence 274931, Sequence 274931,	

ALIGNMENTS

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RESULT 1

US-10-482-433A-3/c

Sequence 3, Application US/10482433A

Publication No. US20040265814A1

GENERAL INFORMATION:
APPLICANT: Epigenomics AG

TITLE OF INVENTION: Method for the detection of cytosine methylation by comparative

TITLE OF INVENTION: Method for the detection of cytosine methylation by comparative

TITLE OF INVENTION: Method for the detection of cytosine methylation by comparative

TITLE OF INVENTION: Method for the detection of cytosine methylation by comparative

TITLE OF INVENTION: Amalysis of single strands of amplificates

FILE REFERENCE: 82174

CURRENT PILING DATE: 2003-12-29

PRIOR APPLICATION NUMBER: US/10/482,433A

CURRENT PILING DATE: 2003-12-29

PRIOR FILING DATE: 2003-12-29

PRIOR APPLICATION NUMBER: DE 10132212.7

PRIOR FILING DATE: 2003-12-29

PRIOR APPLICATION NUMBER: US/10/482,433A

CURRENT Application Product of MdRI-Fragment

US-10-482-433A-3

Query Match

Best Local Similarity 100.0%; Score 24; DB 18; Length 633;

Best Local Similarity 100.0%; Pred. No. 0.22;

Matches 24; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Matches 24; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Matches 24; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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RESULT 4

US-10-451-646-24/c
US-10-451-646-24/c
US-10-451-646-24/c
Sequence 24, Application US/10451646
; Publication No. US20040067508A1
; GENERAL INFORMATION;
GENERAL INFORMATION;
FILTE OF INVENTION: Method for the Simultaneous Amplification of Multiple
TITLE OF INVENTION: Sequences in a PCR Reaction and Marking Thereof
FILE REFERENCE: PCT/DE 01/04951
; FILE REFERENCE: PCT/DE 01/04951
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; OTHER INFORMATION: Amplification Product Of Bisulfite-Treated DNA
US-10-482-433A-6
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; LENGTH: 2932
; TYPE: DNA
; ORGANISM: Homo Sapiens
US-10-473-126-37
            SOFTWARE: PatentIn version 3.1
SEQ ID NO 24
LENGTH: 2932
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/10/482,433A
CURRENT FILING DATE: 2003-12-29
PRIOR APPLICATION NUMBER: DE 10132212.7
PRIOR FILING DATE: 2001-06-27
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin version 3.2
SEQ ID NO 6
TENETH: 633
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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                                                                          CURRENT APPLICATION NUMBER: US/10/451,646
CURRENT FILING DATE: 2003-06-23
NUMBER OF SEQ ID NOS: 25
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Epigenomics AG
TITLE OF INVENTION: Method for the detection of cytosine methylation by comparative
TITLE OF INVENTION: analysis of single strands of amplificates
FILE REFERENCE: 82174
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/10/473,126
CURRENT FILING DATE: 2003-09-26
NUMBER OF SEQ ID NOS: 1258
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TITLE OF INVENTION: Methods and nucleic acids
TITLE OF INVENTION: proliferative disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                               21;
                                                                                                                                                                                                                                                                                                                                                              CTTATATAAACACGTCTTTCAAA 189
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91.3%;
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Pred. No. 23;
0; Mismatches
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Pred. No. 0.28;
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TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement CURRENT APPLICATION NUMBER: US/10/437,963; CURRENT FILING DATE: 2003-05-14

RUMBER OF SEQ ID NOS: 204966

LENGTH: 491

TYPE: DNs
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US-10-473-126-167/c
i Sequence 167, Appl:
Publication No. US:
; OTHER INFORMATION: Clone ID: PAT_MRT4530_85354C.1 US-10-437-963-86300
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CURRENT FILING DATE: 2003-09-26
NUMBER OF SEQ ID NOS: 1258
SEQ ID NO 167
LENGTH: 2932
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Best Local Similarity
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APPLICANT: Epigenomics AG
TITLE OF INVENTION: Methods and nucleic acids
TITLE OF INVENTION: proliferative disorders
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Best Local Similarity
                                 FEATURE:
NAME/KEY: unsure
LOCATION: (1)...(491)
OTHER INFORMATION: unsure at
                                                                                                                                                                                                                                                                                                                                                   APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
                            FEATURE:
                                                                                                        TYPE: DNA
ORGANISM: Oryza sativa
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ORGANISM: Artificial Sequence
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ilarity 91.3%;
Conservative
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; Pred. No. 29;
0; Mismatches
                                           all n locations
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Pred. No. 29;
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APPLICANT: Allen, Keith
APPLICANT: Hoffman, Neil
FIILE OF INVENTION: Expressed Sequences of Arabidopsis
FILE OF INVENTION: Expressed Sequences of Arabidopsis
FILE REFERENCE: 2026 (PARA-015PRV)
CURRENT APPLICATION NUMBER: US/09/770,961
CURRENT FILING DATE: 2001-01-26
PRIOR APPLICATION NUMBER: 60/178,466
PRIOR APPLICATION NUMBER: 60/178,466
PRIOR FILING DATE: 2000-01-27
NUMBER OF SEQ ID NOS: 999
SOFTMARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 577
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US-10-357-930-19156
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; TYPE: DNA
; ORGANISM: Arabidopsis thaliana
US-09-770-961-577
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Best Local
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APPLICANT:
APPLICANT:
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APPLICANT:
APPLICANT:
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              APPLICANT: Schlegel. Robert
APPLICANT: Endege, Wilson
APPLICANT: Endege, Wilson
APPLICANT: Monahan, John
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS FOR
TITLE OF INVENTION: IDENTIFICATION, ASSESSMENT, PREVENTION, AND THERAPY OF
FILE OF INVENTION: HUMAN PROSTATE CANCER
FILE REFERENCE: MRI-007BCN
CURRENT APPLICATION NUMBER: US/10/357,930
CURRENT FILING DATE: 2003-02-04
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PRIOR APPLICATION NUMBER: 09/785,276
                                                                                                                                                                                                                                                                                                                                          297 TTATGTAGACAAGTATGTCAAAG 275
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20; Conservative
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Hamilton, Carol M.
Price, Jennifer L.
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Allen, Keith
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Haas, William David
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Ledford, Brooke L.
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eaka, Joshua G.
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87.0%;
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Pred. No. 1.3e+02;
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PRIOR APPLICATION NUMBER: 60/189,862
PRIOR FILING DATE: 2000-03-16
PRIOR PPLICATION NUMBER: 60/207,454
PRIOR FILING DATE: 2000-05-25
PRIOR APPLICATION NUMBER: 60/211,314
PRIOR FILING DATE: 2000-06-09
PRIOR APPLICATION NUMBER: 60/219,007
PRIOR FILING DATE: 2000-07-18
PRIOR PILING DATE: 2000-07-18
PRIOR APPLICATION NUMBER: 60/255,281
PRIOR FILING DATE: 2000-12-13
NUMBER OF SEQ ID NOS: 62232
SOFTWARE: PASCSEQ for Windows Version 4.0
SEQ ID NO 54995
LENGTH: 597
TYPE: DNA
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, NAME/KEY: misc_feature
; LOCATION: 166, 171
; OTHER INFORMATION: n = A,T,C
US-10-357-930-19156
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US-10-357-930-54995/c
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                                                                                                   ; ORGANISM: Homo sapiens US-10-357-930-54995
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PRIOR APPLICATION NUMBER: 60/207,454
PRIOR FILING DATE: 2000-05-25
PRIOR APPLICATION NUMBER: 60/211,314
PRIOR FILING DATE: 2000-06-09
PRIOR APPLICATION NUMBER: 60/219,007
PRIOR FILING DATE: 2000-07-18
PRIOR FILING DATE: 2000-07-18
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PRIOR FILING DATE: 2000-12-13
NUMBER OF SEQ ID NOS: 62232
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 19156
LENGTH: 520
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 54995, Application US/10357930 Publication No. US20040259086A1 GENERAL INFORMATION:
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Best Local Similarity
Matches 20; Conserv
Query Match 75.8%;
Best Local Similarity 87.0%;
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Schlegel. Robert
APPLICANT: Endege, Wilson
APPLICANT: Monahan, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 2003-02-16 PRIOR APPLICATION NUMBER: 60/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/10/357,930
CURRENT FILING DATE: 2003-02-04
ERIOR APPLICATION NUMBER: 09/785,276
PRIOR FILING DATE: 2003-02-16
PRIOR APPLICATION NUMBER: 60/183,319
PRIOR FILING DATE: 2000-02-17
PRIOR FILING DATE: 2000-02-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: MONAHAN, JOHN
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS FOR
TITLE OF INVENTION: IDENTIFICATION, ASSESSMENT, PREVENTION, AND THERAPY OF
TITLE OF INVENTION: HUMAN PROSTATE CANCER
FILE REFERENCE: MRI-007BCN
FILE REFERENCE: MRI-007BCN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 60/189,862
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  Score 18.2; DB 18;
Pred. No. 1.3e+02;
0; Mismatches 3;
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Pred. No. 1.3e+02;
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                                                  Length 597;
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Gaps

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CTTATGTAGACACGTCTTTCAAA

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Publication ...
GENERAL INFORMATION:
APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Zhou, Yongwei
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-357-930-48960
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SEQ ID NO 86301
LENGTH: 1504
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PRIOR FILLING DATE: 2000-03-16
PRIOR APPLICATION NUMBER: 60/207,454
PRIOR FILLING DATE: 2000-05-25
PRIOR APPLICATION NUMBER: 60/211,314
PRIOR FILLING DATE: 2000-06-09
PRIOR APPLICATION NUMBER: 60/219,007
PRIOR FILLING DATE: 2000-07-18
PRIOR APPLICATION NUMBER: 60/255,281
PRIOR APPLICATION NUMBER: 60/255,281
PRIOR FILLING DATE: 2000-12-13
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Publication No. US20040259086A1
GENERAL INFORMATION:
APPLICANT: Schlegel. Robert
APPLICANT: Endege, Wilson
APPLICANT: Monahan, John
TITLE OF INVENTION: HOVEL GENES, COMPOSITIONS, KITS,
TITLE OF INVENTION: HUMAN PROSTATE CANCER
FILE REFERENCE: MRI-007BCN
GIBERT AND INVENTION: HOMAN PROSTATE CANCER
FILE REFERENCE: MRI-007BCN
                                     TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21(53221)8
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
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CURRENT FILING DATE: 2003-02-04
PRIOR APPLICATION NUMBER: 09/785,276
PRIOR FILING DATE: 2003-02-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 62232
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PRIOR FILING DATE: 2000-02-17
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Wei
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                                                                                                                                                                                                          Boukharov, Andrey A.
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Pred. No. 1.3e
0; Mismatches
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FILE REFERENCE:
CURRENT APPLICATION NUMBER: US/10/473,126
CURRENT FILING DATE: 2003-09-26
NUMBER OF SEQ ID NOS: 1258
SEQ ID NO 168
LENGTH: 2932
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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                                                                                                                         US-10-473-126-313
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US-10-473-126-313/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; ORGANISM: Oryza sativa
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT4530_85355C.1
US-10-437-963-86301
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Publication No. US20040234973A1
GENERAL INFORMATION:
APPLICANT: Epigenomics AG
TITLE OF INVENTION: Methods and nucleic acids
TITLE OF INVENTION: proliferative disorders
                                                             Matches
                                                                                                                                                                                                   SEQ ID NO 313
LENGTH: 2932
                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                           Query Match
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Best Local Similarity 87.0%;
Matches 20; Conservative
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CURRENT APPLICATION NUMBER: US/10/473,126
CURRENT FILING DATE: 2003-09-26
NUMBER OF SEQ ID NOS: 1258
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TITLE OF INVENTION: Methods and nucleic acids for the analysis of hematopoietic cell
TITLE OF INVENTION: proliferative disorders
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                                                                                                                                 OTHER INFORMATION: chemically treated genomic
                                                                                                                                                        ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                        TYPE: DNA
                                                                            Local
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828
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                                                                          Similarity
                  CTTATGTAGACACGTCTTTCAAA 23
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87.0%;
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Pred. No. 1.7e+02;
                                                                       Score 18.2;
Pred. No. 1
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Pred. No. 1
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                                                                                                                                  (Homo sapiens)
                                                                                    Length 2932;
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US-10-027-632-8632 Application US/10027632 Sequence 8632, Application US/10027632 Publication No. US20020198371A1 GENERAL INFORMATION: APPLICANT: Wang, David G. TITLE OF INVENTION: Identification and Mapping of Single Nucleotide TITLE OF INVENTION: Polymorphisms in the Human Genome FILE REFERENCE: 108827.129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 밁
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; ORGANISM: Human
US-10-027-632-8632
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US-10-027-632-8632
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CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR APPLICATION NUMBER: US 60/187,363
PRIOR FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/167,363
                                                                                                                                                                                                                                                        PRIOR PPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR PPLICATION NUMBER: US 60/198,676
PRIOR PRILING DATE: 2000-04-20
PRIOR PRILING DATE: 2000-04-20
PRIOR PRILING DATE: 2000-03-29
PRIOR PPLICATION NUMBER: US 60/193,483
PRIOR PILING DATE: 2000-02-24
PRIOR PILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR PILING DATE: 1999-11-23
PRIOR PILING DATE: 1999-11-23
PRIOR PILING DATE: 1999-09-28
PRIOR PILING DATE: 1999-09-28
PRIOR PILING DATE: 1999-09-28
; TYPE: DNA
; ORGANISM: Human
US-10-027-632-8632
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Best Local Similarity
                                                                                  NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
SEQ ID NO 8632
LENGTH: 619
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Wang, David G.
APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 10887.129
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
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PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR FILING DATE: 1999-09-28
PRIOR APPLICATION NUMBER: US 60/146,002
PRIOR FILING DATE: 1999-08-09
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SOFTWARE: FastSEQ for Windows Version 4.0
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PRIOR APPLICATION NUMBER: US 60/146,002
PRIOR FILING DATE: 1999-08-09
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Minimum DB seq
Maximum DB seq
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Perfect score:
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                           Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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76:
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length: 2000000000
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1: geneseqn1980s:*
2: geneseqn1990s:*
3: geneseqn2000s:*
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SUMMARIES
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Aad39033 Human mdr	Aad39037 Human mdr	Adb92133 Human MDR	Adb96942 Human MDR	Adb87959 Human UGT	Adb20870 MRP1 base	Acf62751 Cancer ba	Aad41243 Human MDR	Abs98184 Human mul	Ads88985 Human ABC	Ade84005 5' regula	Adb53945 MDR1 geno	Abz09897 Human 5'	Human	Human	Aag62614 Human mdr	Abx94595 Human Mdr	Human	Human	Aad39008 Human mdr	Description

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ALIGNMENTS

RESULT 1 AAD39008/c Human; haematologic malignancy; multidrug resistance; MDR; SUMO-1; hypoxia inducible factor-1; small ubiquitin-like-modifier; HIF-1; lymphoid disorder; chronic lymphoproliferative disorder; lymphoma; myeloid disorder; lymphocytic leukaemia; thrombocythaemia; myeloma; angiogenic myeloid metaplasia; myeloid leukaemia; gene therapy; polycythaemia vera; hypoxia responsive element; HRE; ds. Human mdr1 gene HIF-1 binding site DNA #1. AAD39008; AAD39008 standard; DNA; 23-SEP-2002 (first entry) misc_binding Homo sapiens. WO200234291-A2 Location/Qualifiers 10. 14 /*tag= a /bound_moiety= "HIF-1" 24 쁌

02-MAY-2002. 25-OCT-2001; 2001WO-US049856.

(BGHM) BRIGHAM & WOMENS HOSPITAL INC.

26-OCT-2000; 2000US-0243542P.

Colgan SP;

WPI; 2002-471427/50.

Treating a subject (at risk of) having a hematologic malignancy or multidrug resistance, e.g. lymphoma or myeloma, by administering hypoxia inducible factor 1 binding molecules or small ubiquitin-like-modifier-1 binding molecules.

Example

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Page 12; 92pp;

English

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    Treating a subject (at risk of) having a hematologic malignancy or multidrug resistance, e.g. lymphoma or myeloma, by administering hypoxia inducible factor 1 binding molecules or small ubiquitin-like-modifier-1
                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; haematologic malignancy; multidrug resistance; MDR; SUMO-1; hypoxia inducible factor-1; small ubiquitin-like-modifier; HIF-1; lymphoid disorder; hymphoraliferative disorder; hymphoma; myeloid disorder; lymphocytic leukaemia; thrombocythaemia; myeloma; angiogenic myeloid metaplasia; myeloid leukaemia; gene therapy; polycythaemia vera; hypoxia responsive element; HRE; antisense;
                                                                                                                                                                  26-OCT-2000;
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modified_base
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                                                                                                                                     (BGHM ) BRIGHAM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a method of treating a subject having or at risk of developing a haematologic malignancy or multidrug resistance (MDR). The method involves administering hypoxia inducible factor-1 (HIF-1) binding molecules or small ubiquitin-like-modifier (SUMO)-1 binding molecules or small ubiquitin-like-modifier (SUMO)-1 binding molecules or small ubiquitin-like-modifier (SUMO)-1 binding molecules or antisense nucleic acid molecules and SUMO-1 binding molecules or antisense molecules are useful for treating a subject having or at risk of developing haematologic malignancy or MDR (e.g. a lymphoid disorder or a myeloid disorder). The lymphoid disorders include lymphocytic leukaemia or chronic lymphoproliferative disorders e.g. lymphoma, myeloma or chronic lymphoid leukaemia. The myeloid disorders include chronic or acute myeloid leukaemia, e.g. angiogenic invention is used in gene therapy. The present sequence is human mdr1 dene HIF-1 binding cite num
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/note= "Phosphorothioate backbone"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a method of treating a subject having or at risk CC of developing a haematologic malignancy or multidrug resistance (MDR). CC The method involves administering hypoxia inducible factor-1 (HFF-1) CC binding molecules or small ubiquitin-like-modifier (SUMO)-1 binding CC molecules or small ubiquitin-like-modifier (SUMO)-1 binding CC element (HRE) binding molecules or antisense mucleic acid molecules and CC SUMO-1 binding molecules or antisense molecules are useful for treating a CC (e.g. a lymphoid disorder or a myeloid disorder). The lymphoid disorders include lymphocytic leukaemia or chronic lymphoproliferative disorders CC e.g. lymphoma, myeloma or chronic lymphoid leukaemia. The myeloid convention is used in gene therapy. The present sequence is an antisense CC invention is used in gene therapy. The present sequence is an antisense CC oligo targetted to HRE in the mdr1 gene (mdr1-HRE) to inhibit its corporation. This oligo is used in the exemplification of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-NOV-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cancer therapy;
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17-JAN-1995
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This is a preferred fragment of the mdr-1 gene promoter, isolated human DNA. The mdr-1 promoter and enhancer elements are inducible cytostatic agents such as vincristine and adriamycin which are use cancer therapy. Vectors comprising the promoter and enhancer seque

lucible by are used in

from

Claim 2; Page 6; 7pp;

German

cytostatic

agents.

New mammalian expression vector useful for gene therapy, - comprising promoter and/or the enhancer of the mdr-1 gene which is susceptible

comprising the sceptible to

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CC This invention describes a novel method of determining cytosine CC methylation in a sample of genomic DNA which comprises treating the Sample with hydrogen sulphite so that the cytosine is converted to uracil CC whilst 5-methylcytosine remains unchanged, amplifying sections of the DNA CC using at least 2 PCR primers and studying the base composition of both CC complementary amplified strands whereby methylation status is deduced CC from the difference in molecular weight of the two strands. The method is CC used to diagnose and/or prognose unwanted side effects of medication, CC cancer, central nervous system disorders, aggression symptoms or CC behavioural disorders, clinical, psychological and social consequence of CC brain damage, psychotic and personality disorders, dementia and CC associated disorders, cardiovascular disease, malfunction, damage or disease of the gastrointestinal tract, breathing system, bone muscle, CC endocrine or metabolic system, injury, infection, abnormal development or Gexual malfunction. This sequence represents the human Mdzl DNA fragment CC amplified by the PCR primers represented in ABX94593 & ABX94594 and is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   operably linked to heterologous genes coding for therapeutic agents are claimed. Using the vectors, the therapeutic agents (e.g. anti-oncogenes, interferon, IL-2, IL-6, etc.) can be expressed in cancer cells under the control of cytostatic agents. (Updated on 25-MAR-2003 to correct PN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 17; 38pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Determining cytosine methylation in a genomic DNA sample by treating with hydrogen sulfite and analyzing the result, to diagnose associated
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Pred. No.
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                                                                                                                                                                                                                                        This is a preferred fragment of the mdr-1 gene promoter, isolated from human DNA. The mdr-1 promoter and enhancer elements are inducible by cytostatic agents such as vincristine and adriamycin which are used in cancer therapy. Vectors comprising the promoter and enhancer sequences operably linked to heterologous genes coding for therapeutic agents are claimed. Using the vectors, the therapeutic agents (e.g. anti-oncogenes, interferon, IL-2, IL-6, etc.) can be expressed in cancer cells under the control of cytostatic agents. (Updated on 25-MAR-2003 to correct PN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New mammalian expression vector useful for gene therapy, - comprising to promoter and/or the enhancer of the mdr-1 gene which is susceptible to
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17-JAN-1995
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                                                                                                                                                                  Sequence 1318
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   multiple drug resistance; mdr-1; mammalian expression vector;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human mdr-1
                                                                                                                                                                                                                                interferon, II-2, II-5, etc., can control of cytostatic agents. (Updated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-NOV-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          inducible promoter; cytostatic agent; adriamycin; vincristine;
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CTTATGTAGACACGTCTTTCAAAG
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                                                                                                                                                                        374 T; 0 U; 0 Other;
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                                                inducible promoter; cytostatic agent; adriamycin; vincristine;
multiple drug resistance; mdr-1; mammalian expression vector;
cancer therapy; ss.
                                                                                                                                                                                                                                                                                                                                                         Sequence 1688 BP; 556 A; 311 C; 371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 2; Page 5; 7pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cytostatic agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New mammalian expression vector useful for gene therapy, - comprising the promoter and/or the enhancer of the mdr-1 gene which is susceptible to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1994-168680/21.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   inducible promoter; cytostatic agent; adriamycin; vincristine;
multiple drug resistance; mdr-1; mammalian expression vector;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human mdr-1 promoter fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-MAR-2003
17-JAN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ62613;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (DELB-) DELBRUECK CENT MOLEKULARE MEDIZIN MAX
                                                                                                                                                                                                                                                           1323
                                                                                                                                                                                                                                                                                  μ,
                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                         CTTATGTAGACACGTCTTTCAAAG 1300
                                                                                                                                                                                                                                                                        CTTATGTAGACACGTCTTTCAAAG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Walther W;
                                                                                                      promoter fragment.
                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             92DE-04238778
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      92DE-04238778.
                                                                                                                                                                                                                                                                                                                   100.0%; Score 24;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                        G; 450 T;
                                                                                                                                                                                                                                                                                                                   0.14;
                                                                                                                                                                                                                                                                                                                              DB 2;
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                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                               Length 1688;
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                                                                                                                                                                                                                                                                                                         Indels
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Best Local
                                 Berlin K, Braun A, Distler J,
Olek A, Piepenbrock C, Adorjan
Lewin A, Lipscher E, Maier S,
Schwope I, Ziebarth H;
                                                                                                                                                                                                                                  Human; haematopoietic cell proliferation disorder; cytostatic; gene therapy; lymphocytic leukaemia; acute myelogenous leukaem cytosine methylation state; gene; ds.
                                                                                                                                                                                                                                                                                                                                                           9897/c
ABZ09897
                                                                                                                  26-MAR-2001; 2001US-0278333P
                                                                                                                                          26-MAR-2002; 2002WO-EP003401
                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                    ABZ09897;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 2090 BP; 714 A; 365 C; 444 G; 567 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancer therapy. Vectors comprising the promoter and enhancer sequences operably linked to heterologous genes coding for therapeutic agents are claimed. Using the vectors, the therapeutic agents (e.g. anti-oncogenes, interferon, IL-2, IL-6, etc.) can be expressed in cancer cells under the control of cytostatic agents. (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                        WO200277272-A2
                                                                                                                                                                                                                                                                                  Human 5' and/or regulatory region of MDR1 DNA SEQ ID NO:37.
                                                                                                                                                                                                                                                                                                             16-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This is a preferred fragment of the mdr-1 gene promoter, isolated from human DNA. The mdr-1 promoter and enhancer elements are inducible by cytostatic agents such as vincristine and adriamycin which are used in
                                                                                            (EPIG-) EPIGENOMICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 2; Page 4; 7pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New mammalian expression vector promoter and/or the enhancer of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Stein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-NOV-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cytostatic agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-MAY-1994.
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                                                                                                                                                                                                                                                                                                                                                                                                                      1725
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                                                                                                                                                                                                                                                                                                                                                           standard;
                                                                                                                                                                                                                                                                                                                                                                                                                      CTTATGTAGACACGTCTTTCAAAG 1702
                                                                                                                                                                                                                                                                                                                                                                                                                                  CTTATGTAGACACGTCTTTCAAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Walther W;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
ilarity 100.0%;
Conservative (
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                                                                                                                                                                                                                                                                                                                                                          DNA; 2932
                                                                                            AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 24; DB 2;
Pred. No. 0.14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                           Guetig D, P, Grabs Model F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           useful for gene therapy,
the mdr-1 gene which is
                                                                                                                                                                                                                                                                                                                                                                                                                                           24
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                                          G, Lesche R, Leu
Mueller V, Otto T,
                                                                                                                                                                                                                                                  myelogenous leukaemia;
                                                                    Howe A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 2090;
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                                                           Mueller J;
e R. Leu E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              susceptible
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WPI; 2003-018942/01

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RESULT 9
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ADB53945/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            represent specifically claimed nucleotide sequences from the present convention. Oligonucleotides from the present invention can be used: for differentiating between healthy haematopoietic cells and proliferative disorder haematopoietic cells; for differentiating between acute (lymphocytic leukaemia and acute myelogenous leukaemia, as probes for lymphocytic leukaemia and acute myelogenous leukaemia, as probes for determining the cytosine methylation state and/or single nucleotide (polymorphisms (SNPs) of haematopoietic cell proliferation disorder related sequences and their complements; and as primers for the amplification of haematopoietic cell proliferation disorder related DNA sequences. The nucleotide sequences from the present invention can also be used for detecting a predisposition to, differentiation between subclasses, diagnosis, prognosis, treatment and/or monitoring of haematopoietic cell proliferative disorders. The present method enables a highly specific classification of haematopoietic cell proliferative disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       subject. The method comprises contacting a target nucleic acid in a biological sample obtained from the subject with at least 1 reagent which distinguishes between methylated and non-methylated CpG dinucleotides within the target nucleic acid. ABZ09861 to ABZ11118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detecting and differentiating between hematopoietic cell proliferative disorders, comprises contacting a target nucleic acid with a reagent that distinguishes between methylated and non-methylated CpG dinucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    differentiating between haematopoietic cell proliferative disorce associated with at least 1 gene and/or their regulatory regions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 38;
                                                                                                                                                                                  Adorjan P,
Rujan T, 8
                                                                                                                                                                                                                                                                                                                                                                                                 27-FEB-2003; 2003WO-EP002035
                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO2003072821-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      colon cell proliferative disorder; non methylated CpG dinucleotide; cytostatic; cancer; adenoma; carcinoma; cytosine methylation state;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MDR1 genomic DNA region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADB53945;
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Detecting and differentiating between colon cell proliferative disorders associated with a gene or its regulatory regions comprises contacting a target nucleic acid in a biological sample obtained from the subject with
                                                                                                                                                                                                                                                                                                                                     27-FEB-2002; 2002EP-00004551
                                                                                                                                                                                                                                                                             (EPIG-) EPIGENOMICS
                                                                                                                             2003-731620/69
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CTTATGTAGACACGTCTTTCAAAG 805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       allowing for improved and informed treatment of patients
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID
                                                                                                                                                                                  Burger M,
Schmitt A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                invention describes a method for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 841 A; 564 C; 653 G; 874 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                             8
6
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                                                                                                                                                                                                                     Maier S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                        Becker E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 2932;
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         es contacting a the subject with
                                                                                                                                                                                                                        Lesche
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The invention relates to a novel method for detecting and differentiating between colon cell proliferative disorders associated with at least one gene or its regulatory regions. The method comprises contacting a target nucleic acid in a biological sample obtained from the subject with at least one reagent or a series of reagents, where the reagent or series of reagents of aistinguishes between methylated and non methylated CpG dinucleotides within the target nucleic acid. The molecules of the dinvention demonstrate cytostatic activity whilst the method may useful invention demonstrate cytostatic activity whilst the method may useful for detecting and differentiating between colon cell proliferative disorders, including cancers such as colon adenoma and colon carcinoma. The PNA (peptide nucleic acid)-oligomers are useful as probes for determining cytosine methylation state or single nucleotide polymorphisms. The current sequence is that of the genomic DNA region of the invention. This sequence is not shown within the specification but is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 46; SEQ ID NO 1; 74pp; English
                                                          taken from Wipoweb.
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Matches
                                                      Query Match
Best Local
                                                                                      Sequence 2932 BP; 841 A; 564 C; 653 G; 874 T; 0 U; 0 Other;
828 CTTATGTAGACACGTCTTTCAAAG 805
                      -
                                             24;
                                                       Similarity
                     CTTATGTAGACACGTCTTTCAAAG 24
                                             Conservative
                                                         100.0%;
                                              0
                                                       Score 24;
Pred. No.
                                              Mismatches
                                                          0.15;
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                                                                   Length 2932;
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RESULT 10
ADE84005/c
                                                                                                                                                                                                               ds; lymphoid cell proliferative disorder; methylation; methylated CpG dinucleotide; single nucleotide polymorphism; diffuse large B-cell lymphoma; mantle cell lymphoma; chronic lymphocytic leukemia; small lymphocytic lymphoma;
                                                                                                                                                                                                                                                                                                                                          5' regulatory
                                                                                                                                                                                                                                                                                                                                                                                         29-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                           ADE84005;
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25-NOV-2002; 2002WO-EP013265
                                                  30-MAY-2003
                                                                                               WO2003044226-A2
                                                                                                                                            Homo sapiens
                                                                                                                                                                                              follicular lymphoma; diagnosis; prognosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard; DNA; 2932
                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                          region of human gene MDR1.
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23-NOV-2001; 28-DEC-2001;

2001DE-01057491. 2001DE-01064501.

Burger M,

Caldwell

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Genc

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Becker E,

Maier S,

Nimmrich

2003-457621/43.

(EPIG-) EPIGENOMICS

Detecting and differentiating between lymphoid cell proliferative disorders comprises contacting a target nucleic acid with at least one reagent that distinguishes between methylated and non-methylated CpG

The invention relates to a method of detecting and differentiating between lymphoid cell proliferative disorders associated with at least

SEQ

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448pp; English

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RESULT 11
ADS88985/c
ID ADS889
XX ADS889
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XX IS Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 24
                                                                                                                                                              Predicting responsiveness of a subject with breast cell disorder, useful for treating or differentiating breast proliferative disorders comprises analyzing methylation genomic DNA from the subject.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Foekens J,
Nimmrich I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-OCT-2002; 2002DE-01045779
07-JAN-2003; 2003DE-01000096
17-APR-2003; 2003DE-01017955
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-OCT-2003; 2003WO-EP010881
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human ABCB1 gene SEQ ID. NO:1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADS88985;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADS88985 standard; DNA; 2932
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (EPIG-) EPIGENOMICS AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 2932 BP; 841 A; 564 C;
                                                                                                                                                                                                                                                                                                                                                             2004-348468/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24;
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Harbeck N,
Rujan T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cell proliferative disorder; breast; methylation; therapy; single nucleotide polymorphism; SNP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                          Koenig
Schmitt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
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Pred. No. 0.15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Maier S, N
Schmitt M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 10;
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1, Look N
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The invention relates to

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novel method

for

predicting

the

responsiveness

Isolated nucleic acid molecules having polymorphisms in known human ge e.g. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for

Example 2;

SEQ ID

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1; 104pp; English.

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RESULT 12
ABS98184/c
ID ABS981
XX Human;
XX Human;
XX Cytoch
XX Cytoch
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XX Gytoch
XX Human;
XX Human;
XX Human;
XX Human;
XX Gytoch
XX Gy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        KW Human; ds; cytochrome P450 A1; CYP4501A1; UGT2B4; MDR1;

KW cytochrome P450 A2; CYP4501A2; cytochrome P450 02E; CYP45002E1; LTF;

KW adrenergic receptor beta1; ADBR1; aryl hydrocarbon; AHR; MRP3; NR112;

KW aryl hydrocarbon receptor nuclear translocator; ARNT; cathepsin S; CTSS;

KW cyclooxgenase 2; COX2; diazepam binding inhibitor; DBI; haematological;

KW cyclooxgenase 2; EPHX2; 5-lipoxygenase activating protein; FLAP;

Byltathione-5-transferase 12; GST12; histamine-N-methyl transferase;

KW HNMT; kallikrein 2; KLK2; nicotinamide-N-methyl transferase; NNNT;

KW NADPH quinone oxidoreductase 21; NQO2; sulfotransferase thermolabile; STM;

KW WHOP-glucuronosyl transferase 224; UDD-glucuronosyl transferase 227;

KW UDP-glucuronosyl transferase; UGT2B15; urokinase receptor; uPA;

KW UGT2B7; UDP-glucuronosyl transferase;

KW multidrug resistance 1; lactotransferin; orphan nuclear receptor;

KW multidrug resistance associated protein 3; cancer; prostate;

MC multidrug resistance associated protein 3; cancer; prostate;

MC multidrug resistance associated protein; colorectal tumour;

KW actylcholine muscarinic receptor; cUMR1; CHMR2; CHMR3; CHMR4; CHMR5;

KW actylcholine muscarinic receptor; pulmonary; immunological; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  밁
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Best Local :
                                                                                                                                          WPI; 2002-698522/75
                                                                                                                                                                                                                                                                                                                                                      28-NOV-2000; 2000US-00724389
                                                                                                                                                                                                                                                                                                                                                                                                                    28-NOV-2001; 2001WO-US044838
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      single nucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200257410-A2
                                                                                                                                                                                                                                                                                     (DNAS-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human multidrug resistance gene promoter B and exons 1-3 sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-DEC-2002
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                                                                                                                                                                                                                                                                                 DNA SCI LAB
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                                                                                                                                                                                                               Hall
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Pred. No.
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XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     밁
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention relates to the sequence of an isolated nucleic acid molecule comprising at least one base variation from that of a known human cytochrome P450 A1 (CYP4501A1), cytochrome P450 A2 (CYP4501A2), cytochrome P450 02E1 (CYP45002E1), adrenargic receptor beta1 (ADBR1),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    disorder-related traits.
                                                                                                                                                                                                                                                      Human; tranagenic; reporter construct; CYP3A4; cytochrome P450; transcriptional regulatory element; xenobiotic; steroid; MDR1; p-glycoprotein; gene; ds.
                                                                                                                                                                                                                                                                                                                                                                                                          Human MDR1 gene 5' flanking region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAD41243 standard; DNA; 10200 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 8573 BP;
                                                 10-MAY-2002
                                                                                                                     WO200236784-A1
                                                                                                                                                                                      Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CTTATGTAGACACGTCTTTCAAAG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 24; DB 6;
Pred. No. 0.16;
Mismatches 0
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RESULT 14
ACF6751/c
ID ACF62751 standard; DNA; 177380 BP
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to generation of non-human transgenic animals comprising a reporter construct for producing a detectable amount of a reporter molecule operably linked to a transcriptional regulatory nucleic acid molecule from the human CYP3A4 gene. The invention also relates to transgenic animal for analysing CYP3A4 cytochrome P450 gene regulation. The non-human mammal of the invention is useful for determining whether a compound is capable of effecting the transcription of a human CYP3A4 gene. It is also useful for determining the effect of a compound such as a xenobiotic ox steroid on the regulation of expression of the CYP3A4 gene in a human. The present sequence is 5' flanking region of human (p-glycoprotein) MDR1 gene. This sequence is used as a regulatory element in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 10200 BP; 3296 A; 1785 C; 1985 G; 3134 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Non-human mammal for determining the effect of a compound on regulation of CYP3A4 gene expression, comprises a reporter construct for producing reporter linked to a transcriptional regulatory nucleic acid from a hum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-537301/57,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-NOV-2000; 2000AU-00001161.
10-MAY-2001; 2001AU-00004901.
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                                                                                                                                                                                                                                                                         Cancer; CYP3A5; irinotecan; pharmaceutical; malignant glioma; cytochrome p450; subfamily IIIA; nifedipine oxidase; polypeptide cytostatic; gene; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 11; Page 83-92; 95pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Liddle C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (UNSY ) UNIV SYDNEY.
                                                                                                                                                                                                                                                                                                                                        Cancer based on CYP3A5 related polynucleotide SEQ ID NO:683.
                                                                                                                                                                                                                                                                                                                                                                        08-OCT-2003 (first entry)
                                                                                                        23-JUL-2001; 2001EP-00117608.
24-MAY-2002; 2002EP-00011710.
                                                                                                                                                       23-JUL-2002; 2002WO-EP008219
                                                                                                                                                                                    20-FEB-2003.
                                                                                                                                                                                                                                               Unidentified
                                                                                                                                                                                                                 WO2003013534-A2
                                                                          (EPID-) EPIDAUROS BIOTECHNOLOGIE AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9051 CTTATGTAGACACGTCTTTCAAAG 9028
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                             Kerb R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; S
100.0%; F
tive 0;
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WPI; 2003-268144/26

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RESULT 15
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XX ADB208
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KW Irinot
KW Iung c
KW Varian
KW Varian
KW Varian
KW V2003
XX V02003
XX V02003
XX EPID-
XX BARNER
PF 23-JUL
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XX Y
PA (EPID-
XX Heinri
XX Disclc
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PF Use of
PF Variar
PF Variar
PF Variar
XX Disclc
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Best Local S
Matches 24
                                                                                                                                               Use of irinotecan or its derivative for preparation of a pharmaceutical composition for treating cancer in a subject having a genome with a variant allele comprising a multidrug resistance protein 1 polynucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             irinotecan; colorectal cancer; cervical cancer; gastric cancer;
lung cancer; ovarian cancer; pancreatic cancer; malignant glioma;
variant allele; multidrug resistance protein 1; MRP1; cytostatic; gene;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-JUL-2001; 2001EP-00117608.
24-MAY-2002; 2002EP-00011710.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (EPID-) EPIDAUROS BIOTECHNOLOGIE
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Pred. No. 0.23;
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The present

invention describes a method for

the use of irinotecan

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Disclosure;

SEQ

ID NO

683; 100pp; English.

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its derivative for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a multidrug resistance protein 1 (MRPI) polynucleotide (II). (I) has cytostatic activity. (I) or its derivative can be used for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject, where the subject is a human (preferably African or Assian) or a mouse. The present sequence represents a sequence which is used in the exemplification of the present invention.
   Sequence 177380 BP; 58338 A; 32530 C; 31389 G;
55123 T; 0 U; 0 Other;
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Matches Query Match Best Local Similarity _ 24; Conservative 100.0%; 0; Score 24; [Pred. No. 0. Mismatches DB . 23; 8 0; Length 177380; Indels 0, Gaps

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멼 Ś 140580 CTTATGTAGACACGTCTTTCAAAG 140557 CTTATGTAGACACGTCTTTCAAAG 24

Search completed: February Job time: 237.862 secs 9 2005, 16:05:52

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Result
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Maximum Match 100%
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 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Match
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24
1 cttatgtagacacgtctttcaaag
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AX504306 Sequence
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	82.5	82.5	82.5	82.5	82.5	82.5	82.5	82.5	82.5	82.5	82.5	82.5	83.3		٠,	•	87.5	87.5	91.7	91.7	95.8	95.8	100.0	100.0	100.0	100.0	
	174986	브	110000										20	158344	110000	21	21	21	22	22	23	23	177380	177380	177380	2932	
	5	N	N	σ	σ	6	σ	σ	6	თ	0	σ	δ	N	N	0	9	σ	თ	σ	σ	σ	9	თ	σ	9	,
ALIGNMENTS	AL935126	CR354542	AC130739_1	AX825881	AX822241	AX795733	AX767413	AX598827	AX529569	CQ806797	AX457065	AX701744	AX504355	CR352257	AC110967_2	AX504354	AX504351	AX504350	AX504353	AX504349	AX504352	AX504348	AC002457	AX707915	AX706985	HSMDR1A	
	AL935126 Zebraiish		tion (2 o					Sequenc	Sequenc	Sequenc					tion (3 o	AX504354 Sequence								AX/0/915 Sequence		_	

RESULT 2 AX504323/c LOCUS LOCUS DEFINITION ACCESSION VERSION VERSION KEYWORDS SOURCE ORGANISM	Qy da	Query Match Best Local Matches 2	source ORIGIN	JOURNAL FEATURES	REFERENCE AUTHORS TITLE	KEYWORDS SOURCE ORGANISM	RESULT 1 AX504306 LOCUS DEFINITION ACCESSION VERSION
AX504323 24 bp DNA linear PJ Sequence 26 from Patent WO0234291. AX504323 AX504323.1 GI:23386135 Homo sapiens (human)	1 CTTATGTAGACACGTCTTTCAAAG 24 	100.0%; Score 24; DB 6; Length 24; Similarity 100.0%; Pred. No. 2.2; 4; Conservative 0; Mismatches 0; Indels		THE BRIGHAM AND WOMEN'S HOSPITAL, INC. (US) LOCATION/Qualifiers	for treating hematologic	Homo sapiens (human) Homo sapiens Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.	AX504306 24 bp DNA linear PAT Sequence 9 from Patent WO0234291. AX504306 AX504306.1 GI:23386124
PAT 27-S		0; Gaps			malignancies	Euteleostomi;	
27-SEP-2002		s 0;			es and	comi;	27-SEP-2002

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REFERENCE
AUTHORS
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AR080195/c
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                                                                                                                                                                                                                                                                                                               203
                                                               Distler, J. and Leu, E.

Method for detecting cytosine methylation analysing single strands of amplificates patent: WO 03002760-A 3 09-JAN-2003;
                                                                                                                                  synthetic construct
synthetic construct
other sequences; artificial sequences.
                                                                                                                                                                                                              AX701741 633 bp
Sequence 3 from Patent WO03002760.
AX701741
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                                                          Epigenomics AG (DE)
                                                                                                                                                                                                  AX701741.1
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Stein, U. and Walther, W.
Vector for the expression of therapy-relevant
Patent: US 5968735-A 4 19-OCT-1999;
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Patent: WO 0234991-A 26 02-MAY-2002;
THE BRIGHAM AND WOMEN'S HOSPITAL, INC.
Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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organism="synthetic construct"
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                      1 CTTATGTAGACACGTCTTTCAAAG 24
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Chen, C.J., Clark, D., Ueda, K.,
Roninson, I.B.
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M29423 J05168 M18754
M29423.1 GI:187472
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Isolation and sequence of the promoter region
multidrug-resistance (P-glycoprotein) gene
J. Biol. Chem. 262 (36), 17432-17436 (1987)
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                                                              Similarity
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                       /organism="Homo sapiens"
/mol_type="genomic DNA"
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/map="'g21"
t<1...>976
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/note="PGY1 m
<1. .244
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/note="PGY1 mRNA
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GDB: GDB:583426
GDB DSEG: PGY1
CONTRACT: Eric D. Green
Genome Technology Branch
National Human Genome Research Institute/NIH
49 CONVENT Dr., MSC4431, Bldg. 49, Rm. 2A08,
Tel: 3014020201
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Bouffard,G.G., Iyer,L.M., Idol,J.R., Braden,V.V., Cunningham
Bouffard,G.G., Mohr-Tidwell,R.M., Peluso,D.C., Fulton,R.S.,
Weintraub,L.A., Mohr-Tidwell,R.M., Peluso,D.C., Fulton,R.S.,
Leckie,M.P. and Green,E.D.
A collection of 1814 human chromosome 7-specific STSs
Genome Res. 7 (1), 59-64 (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: egreen@nhgri.nih.gov
Primer A: AACGGAAGCCAGAACATTCC
Primer B: AGGCTTCCTGTGGCAAAGAG
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Unpublished (1997)
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                                                                                                                                     This STS was developed from sequence determined by another investigator. See GenBank record: M29423 For additional information about the NGRI chromosome 7 mapping project, see http://www.nhgri.nih.gov/DIR/GTB/CHR7. Also see Genomics 11:548-64 (1991) [MUID-92128937].
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Polymerization:
PCR Cycles:
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/gene="PGY1"
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Best Local Similarity 100.0%;
Matches 24; Conservative 0
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Stein,U. and Walther,W.
Stein, U. and Walther,W.
Sector for the expression of therapy-relevant genes
Patent: US 5968735-A 3 19-OCT-1999;
Location/Qualifiers
1. .1318
                                                                 Original
4a) bone
                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 1327)

Madden,M.J., Morrow,C.S., Nakagawa,M., Goldsmith,M.E.,
Fairchild,C.R. and Cowan,K.H.

Identification of 5' and 3' sequences involved in the regulation of transcription of the human mdrl gene in vivo

J. Biol. Chem. 268 (11), 8290-8297 (1993)
                                                                                                                                                                                                                                     L07624:1 GI:187500
MDR1 gene; P-glycoprotein; multidrug resistance protein.
Homo sapiens (human)
                                                                                                                                                                                                                                                                                                  HUMMDR1P 1327 bp DNA
Human multidrug resistant P glycoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unknown
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                                                                                                                                                                                                                         Homo sapiens
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253. .272
/gene="PGY1"
complement(413.
                                                              marrow DNA.
/mol_type="genomic DN/db_xref="taxon:9606"
                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mol_type="unassigned
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                        organism="Homo sapiens"/
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patent US 5968735.
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Pred. No. 1.1;
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                                                                     Stein, U. and Walther, W.
Vector for the expression of therapy-relevant genes
Patent: US 5968735-A 2 19-OCT-1999;
Location/Qualifiers
1. 1688
                                                                                                                                                                                                             1688 bp
Sequence 2 from patent US 5968735.
AR080193
                                                                                                                                                                Unknown.
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                                                                                                                                       (bases 1 to 1688)
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                                              /organism="unknown"
/mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note="major transcription 1041. .1170 /gene="PGY1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /tissue type="bone marrow"
/tissue_lib="lambda Charon 4a"
863. .1171
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/gene="PGY1"
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function="heat shock element"
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bound moiety="unknown"
function="heat shock element"
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A38669/c
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                                                                                                                                                                                                               Eukaryoča; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 2117)

Stein, U. and Walther, W.

VECTOR FOR THE EXPRESSION OF THERAPY-RELEVANT GENES
Patent: WO 9411522-A 1 26-MAY-1994;
MAX DELBRUECK CT FUER MOLEKULA (DE)

Other publication DE 4238778 940519.

Location/Qualifiers
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Sequence
AR080192
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Sequence 1 from Patent WO9411522.
A38669
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Stein,U. and Walther,W.
Vector for the expression of therapy-relevant genes
Patent: US 5968735-A 1 19-OCT-1999;
Location/Qualifiers
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l Similarity 100.0%;
24; Conservative 0;
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                                                                       /replace="t"
                                                                                                replace="c"
                                                                                                                        cell_type="SARCOMAS"
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                                                                                                                                                   /mol_type="unassigned
/db_xref="taxon:9606"
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from patent US 5968735.
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100.0%; Pred. No. 0.96;
tive 0; Mismatches
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Pred. No. 0.96;
Mismatches 0
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Matches 2
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M57450.1 GI:187470
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Human multidrug-resistance (MDR1)
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Kohno,K., Sato,S., Uchiumi,T., Takano,H., Kato,S. and Kuwano,M.
Tissue-specific enhancer of the human multidrug-resistance (MDR1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 multidrug resistance.
                                                                                  Sequence 1 from Patent CQ806551 GI:47111933
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     J. Biol. Chem.
91060578
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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                                               Homo sapiens (human)
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                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
Foekens,J.,
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951._.955
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Harbeck, N.,
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Berlin,K., Braun,A., Distler,J., Guetig,D., Howe,A., Mueller,J., Olek,A., Piepenbrock,C., Adorjan,P., Grabs,G., Lesche,R., Leu,E., Lewin,A., Lipscher,E., Maier,S., Model,F., Mueller,V., Otto,T., Pelet,C. and Ziebarth,H. Methods and nucleic acids for the analysis of hematopoietic cell proliferative disorders
Patent: WO 02077272-A 37 03-OCT-2002;
                                                                                                               Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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source /or /mo /st /db /cl /db /cl /la a /cl /mo /mo /mo /or /mo	CANADA Tel: (613) 755 Fax: (613) 755 Fax: (613) 755 Email: harrisl FEATURES	TITLE Expressed Sequence Tag stage perithecia JOURNAL Unpublished (2004) Contact: Harris, Linds Eastern Cereal and Oil Agriculture and Agri-	ACCESSION CN813365 VERRION CN813365.1 GI:4783737 KEYWORDS EST. SOURCE Gibberella zeae (aname organism) Fungi; Asco Hypocreomycetidae; Hyp (bases 1 to 213) REFERENCE 1 (bases 1 to 213) AUTHORS COuroux, P., De Moors, J. Robert, J. S., Singh, J. J.	RESULT 1 CN813365/C LOCUS CN813365 CN813365 DEFINITION F906 06110 R F906 AAF		17.8 74.2 555 17.8 74.2 559 17.8 74.2 585	17.8 74.2 516 17.8 74.2 522 17.8 74.2 542	C 36 18.2 75.8 986 9 C 37 18.2 75.8 174 6 38 18 75.0 512 8 39 18 75.0 569 8	32 18.2 75.8 749 33 18.2 75.8 75.8 34 18.2 75.8 811 35 18.2 75.8 975	28 18.2 75.8 691 29 18.2 75.8 691 30 18.2 75.8 720 31 18.2 75.8 737	18.2 75.8 652 18.2 75.8 672 18.2 75.8 685

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/organism="Gibberella zeae" /mol_type="mRNA" /mol_type="mRNA" /strain="DAOM 180378" /db xref="taxon:5518" /db xref="taxon:5518" /dev stage="Sexual" /lab host="E. coli DH10B" /lab host="E. coli DH10B" /lab host="E. coli DH10B" /note="Vector: pGem-T easy; Site 1: EcoRI; Mycelia grown on carrot agar at 20oC until confluent; perithecia induced with Tween 40 solution (25% v/v). Fruiting bodies were collected 20 days after induction. Total RNA was extracted using Trizol. cDNAs were amplified using Invitrogen GeneRacer kit. cDNA was not fractionated and was bidirectionally cloned."	Eastern Cereal and Oilseed Research Centre Agriculture and Agri-food Canada Bldg. 21, Central Experimental Farm, Ottawa, Ontario, KIA 0C6, CANADA Tel: (613) 759-1314 Fax: (613) 759-6566 Email: harrislj@agr.gc.ca. Location/Qualifiers 1213	Lacroix,C., Tinker,N.A. graminearum	CN81365. CN81365.1 GI:47837376 EST. Gibberella zeae (anamorph: Fusarium graminearum) Gibberella zeae Gibberella zeae Eukaryota; Fungi; Ascomycota; Pezizomycotina; Sordariomycetes; Hypocreomycetidae; Hypocreales; Nectriaceae; Gibberella. 1 (bases 1 to 213) Harris.L.J., Rocheleau, H., Ouellet, T., Allard, S., Chapados, J., Harris.L.J., Rocheleau, H.,	CN813365 213 bp mRNA linear EST 01-JUN-2004 Fg06_06h10_R Fg06_AAFC_ECORC_Fusarium_graminearum_perithecia Gibberella zeae cDNA clone Fg06_06h10, mRNA sequence.

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Computer-based methods for the mouse full-length cDNA control contro
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Please visit our web site (http://genome.gsc.riken.go.jp) for further details.
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Carninci, P., Shibata, Y., Hayatsu, N., Sugahara, Y., Shibata, K.,
Itoh, M., Konno, H., Okazaki, Y., Muramatsu, M. and Hayashizaki, Y.
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prepare full-length cDNA libraries for rapid discovery of new
genes. Genome Res. 10 (10), 1617-1630 (2000)
wagi, K., Fujiwake, S., Inoue, K., Togawa, Y., Izawa, M., Ohara, E.,
Watahiki, M., Yoneda, Y., Ishikawa, T., Ozawa, K., Tanaka, T.,
Matsuura, S., Kawai, J., Okazaki, Y., Muramatsu, M., Inoue, Y., Kira, A.
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Arakawa,T., Carninci,P., Fukuda,S., Furuno,M., Hanagaki,T.,
Hara,A., Hiramoto,K., Hori,F., Ishii,Y., Ito,M., Kawai,J.,
Konno,H., Kouda,M., Koya,S., Matsuyama,T., Miyazaki,A., Nomura,K.,
Ohno,M., Okazaki,Y., Okido,T., Saito,R., Sakai,C., Sakai,K.,
Sano,H., Sasaki,D., Shibate,K., Shinagawa,A., Shiraki,T.,
Sogabe,Y., Suzuki,H., Tagami,M., Tagawa,A., Shiraki,T.,
Takeda,Y., Tanaka,T., Toya,T., Muramatsu,M. and Hayashizaki,Y.
Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Laboratory for Genome Exploration Research Group, RIKEN Genomic Sciences Center(GSC), Yokohama Institute
The Institute of Physical and Chemical Research (RIKEN)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           u
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RIXEN integrated sequence analysis (RISA) system--384-format sequencing pipeline with 384 multicapillary sequencer. Genome Res. 10 (11), 1757-1771 (2000)

Konno,H., Fukunishi,Y., Shibata,K., Itoh,M., Carninci,P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        and Hayashizaki,Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan
Tel: 81-45-503-9222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Yoshihide Hayashizaki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BB662530
BB662530.1 GI:16496284
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TATGTAGACACGGCTTTCAAAG 165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TATGTAGACACGTCTTTCAAAG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   81-45-503-9216
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RIKEN full-length enriched, 15 days embs
cDNA clone D930005N11 5', mRNA sequence
               /tissue_type="head"
                                                                                                                           /mol_type="mRNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                              1. .671
                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                       'sex="mixed"
                                                                                           clone="D930005N11"
                                                                                                                                                                                                                                                   organism="Mus musculus"
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95.5%;
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Pred. No. 50;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
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BZ984511
LOCUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURES
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KEYWORDS
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Query Match
Best Local Similarity
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Query Match
Best Local S
Matches 21
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BZ984511
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                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Viridiplantae; Streptophyta; En
Spermatophyta; Magnoliophyta; Liliopsida;
clade; Panicoideae; Andropogoneae; Zea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                Zea mays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BZ984511.1
GSS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 TTATGTAGACACGTCTTTCAAAG 24
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                                                                                                                                                               Class: sheared ends.
                                                                                                                                                                              seq primer: TF
                                                                                                                                                                                                                                                                        Contact: Cathy Whitelaw
                                                                                                                                                                                                                                                                                                      Unpublished (2003)
                                                                                                                                                                                                                                                                                                                     Maize Genomics Consortium
                                                                                                                                                                                                                                                                                                                                       Bennetzen, J
                                                                                                                                                                                                                                                                                                                                                      Resnick, A.,
                                                                                                                                                                                                                                                                                                                                                                    Whitelaw, C.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PUGIE79TD ZM_0.6_1.0_KB Zea genomic survey sequence.
                                                                                                                                                                                           Email: whitelaw@tigr.org
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Zea mays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BZ984511
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itelaw, C.A., Quackenbush, J.,
                                                                                                                                                                                                           301-838-5843
301-838-0208
                                                                                                                                                                                                                                         Medical Center Drive,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone="ZMMBTa390M13"
/clone_lib="ZM_0.6_1.0 KB"
/note="Vector: pCR4-TOPO; Site_1:
CoT selected genomic DNA library"
                                                                                  organism="Zea mays"
/mol_type="genomic DNA"
/strain="B73"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note="Site 1: Sall; Site 2: BamHI; cDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Riken Genomic Sciences Center and Genome Science Laboratory in RIKEN, Division of Experimental Animal Research in Riken
                                                                 db_xref="taxon:4577"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone_lib="RIKEN full-length enriched, 15 days embryo
head"
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                                                                                                                                                                                                                                                                                                                                                  Fraser, C.M.,
                                                                                                                                 . 780
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GI:29219382
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Pred: No. 1.1e+02;
0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                    Yuan,Y.,
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                                                                                                                                                                                                                                         Rockville,
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                                                                                                                                                                                                                                                                                                                                                    San
                                                                                                                                                                                                                                                                                                                                                 Aken,S., Utterback,T., San Miguel,P., Ma,J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA
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                                                                                                                                                                                                                                                                                                                                                                                                            Embryophyta; Tracheophyta;
a; Poales; Poaceae; PACCAD
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                   EcoRI; 0.6-1.0
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                   õ
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80.0**%;** 87.5**%**;

Score 19.2; DB 8; Pred. No. 2.3e+02;

Length 780;

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REFERENCE
AUTHORS
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KEYWORDS
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CC365078/c
                                                                                                                                                                                                                             RESULT 5
BZ984507/c
                          REFERENCE
AUTHORS
                                                                                                                                                    ACCESSION
VERSION
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                                                                                                        ORGANISM
                                                                                                                                       KEYWORDS
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                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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                                                                                                                                                                                                                                                                                                 103
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CC365078
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1 (bases 1 to 780)

Whitelaw, C.A., Quackenbush, J., Van Aken, S., Utterback, T.,

Whitelaw, C.A., Quackenbush, J., Van Miguel, P., Ma, J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD
clade; Panicoideae; Andropogoneae; Zea.
                                                                                                                                                                                BZ984507

PUGIE79TB ZM_0.6_1.0_KB
genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                              21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seq primer: TR
Class: sheared ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Maize Genomics Consortium Unpublished (2003) Other GSSs: PUHME62TD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Zea mays
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        1 (bases 1 to 816)
Whitelaw,C.A., Quackenbush,J., Van
Resnick,A., Fraser,C.M., Yuan,Y.,
                                                      Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD clade; Panicoideae; Andropogoneae; Zea.
                                                                                                                                         GSS
                                                                                                                                                    BZ984507
BZ984507.1 GI:29219374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: whitelaw@tigr.org
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Cathy Whitelaw
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/clone="1b="ZM_0.6_1.0 KB"
/note="Vector: pCR4-TOPO; Site 1: EcoRI; 0.6-1.0
/note="Vector: pCR4-TOPO; DNA library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      organism="Zea mays"
/mol_type="genomic DNA"
/strain="B73"
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Pred. No. 2.3e+02;
0; Mismatches 3;
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           Van Aken,S., Utterback,T.,
Y., San Miguel,P., Ma,J. and
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Other_GSSs: PUGIE79TD
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Other_GSSB: PUIKG34TBB
Contact: Cathy Whitelaw
                                                                                                                                                                                                                                                                                                                                                                                                  1 (bases 1 to 889)
Whitelaw, C.A., Quackenbush, J., Van Aken, S., Utterback, T.,
Resnick, A., Fraser, C.M., Yuan, Y., San Miguel, P., Ma, J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                               Zea mays
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD
clade; Panicoideae; Andropogoneae; Zea.
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Email: whitelaw@tigr.org
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Fax: 301-838-0208
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/clone=lib="ZM 0.6.1.0 KB"
/clone_lib="ZM 0.6.1.0 KB"
/note="Vector; pCR4_TOPO; Site_1; EcoRI; 0.6-1.0 kb
CoT selected genomic DNA library"
                                          /clone="ZMMBTa0602F19"
/clone lib="ZM_0.6_1.0 KB"
/note="Vector: pCR4-TOPO; Site_1:
CoT selected genomic DNA library"
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/strain="873"
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/mol_type="genomic D
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/mol_type="genomic DNA"
/strain="B73"
/db_xref="taxon:4577"
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  Eschscholzia californica (California poppy)
Eschscholzia californica
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; Ranunculales;
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.

1 (bases 1 to 426)

1 (bases 1 to 426)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (2004)
Contact: Rod A. Wing
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OR_BBa0082M07.r OR_BBa Oryza
3', genomic survey_____
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CL745575.1 GI:50686923
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/note="Vector: pAGIBAC1; Site_1: HindIII; Site_2: HindIII"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    tissue_type="young leaves"
/lab_host="DH10B-T1 phage resistant"
/clone_lib="OR_BBa"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /mol_type="genomic DNA"
/db_xref="taxon:4529"
/clone="OR_BBa0082M07"
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BH332334
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                                                                                                                                                                                                                                                                                                         332 TATGTAGGCACGTCTTTCATAG
                  Rattus norvegicus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                     BH332334 627 bp
CH230-125M16.TV CHORI-230 Segment
                                                                                                                    GSS
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                                                                                                                                                      BH332334
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                                                                                          Rattus norvegicus (Norway rat)
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                                                                                                                                                                    CH230-125M16,
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208 Mueller Laboratory, Department of Biology, A'
State University, University Park, PA 16802, USA
Tel: 814 863 6413
Fax: 814 865 9131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: cwd3@psu.edu or jhl10@psu.edu
The sequence provided is trimmed of vector and low quality regions.
Full sequence and original trace file are available from the Plant
Genome Network website (http://pgn.cornell.edu)
Plate: eca01.18ms1 row: f column: 06
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depamphilis,C., Carlson,J., Ma,H., Tanksley,S., Field,D.,
depamphilis,C., Carlson,J., Zahn,L.M., Kong,H., Ilut,D
Druckenmaniler,M., Landherr,L., Hu,Y., Plock,S., Wall,K.,
Chiorean,S., Albert,V., Doyle,J., Frohlich,M., Miller,W.,
Oppenheimer,D., Soltis,D., Soltis,D. and Theisean,G.
Generation of ESTs from early flower buds of Eschscholzia
                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (2002)
Contact: Claude dePamphilis or James Leebens-Mack
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mueller Laboratory
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(bases 1 to 627)
                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /cione lib="Ecaol"
//cione lib="Ecaol"
//note="Vector: pBluescript SK (+/-); Site 1: EcoRI;
Site 2: Xho1; Plants were grown in greenhouse at Penn
State from commercially available seeds. Only floral buds
with diameter of 2.5 mm of less were collected. This is a
directionally cloned, non-normalized library. Avg. insert
length: 1702; Primers: M13F and M13R; Antibiotic: 50 ug/ml
Ampicillin; Primary Titer: 7E6 pfu total; Amplified Titer:
1.68E11 pfu/ml; Mass Excised Titer: 5.68E total; This
library has been generated by the Floral Genome Project
(FGP). We would like to thank Huck Life Sciences
Consortium for their assistance. The Floral Genome Project
is funded by NSB's Plant Genome Research Program
(DBI-0115684). More information about the project can be
obtained at http://fgp.bio.psu.edu"
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/clone="eca01-18ms1-f06"
/tlssue_type="flower buds <= 2.5mm"</pre>
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/lab_host="SOLR"
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Pred. No. 3.3e+02;
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sequence.~
                                                                                                                                                                                                                                                                                                                                                                                                                    Length 464;
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Clones are derived from the rat BAC library CHORI-230
Clones are derived from the rat BAC library CHORI-230
(http://www.chori.org/bacpac/rat230.htm). For BAC library
(http://www.chori.org/bacpac/credition (pdejong@mail.cho.org).
Clones may be purchased from BACPAC Resources
(http://www.chori.org/bacpac/or ering_information.htm). BAC end
page: http://www.tigr.org/tdb/bac_ends/rat/bac_end_intro.html
Plate: 125 row: M column: 16
Seq primer: T7
Class: BAC ends.
                                                                                                                                                                                                                                                                                                                                                                                                                  OR_BBa0024M03.r OR_BBa Oryza
3', genomic survey sequence.
CL706910
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                                                                                                                                                                                                                Oryza rufipogon
Cuyas rufipogon
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
1 (bases 1 to 632)
Kim,H., Yu,Y., Stum,D., Yost,D., Rao,K., Luo,M., Jetty,R.,
Kudrna,D., Muller,C., Hatfield,J., Soderlund,C. and Wing,R.
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20; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Other_GSSs: CH230-125M16.TJ
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Rat BAC End Sequences from Library CHORI-230 EcoRI segment
                           University of Arizona
Forbes Building Room 303, Tucson,
Tel: 520 626 5959
Fax: 520 621 1259
Email http://genome.arizona.edu
                                                                                                                                                                            OMAP Project
Unpublished (2004)
                                                                                                                                                                                                                                                                                                                                                           Oryza rufipogon
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                                                                                                                                      Contact: Rod A. Wing
Arizona Genomics Institute
                  PCR PRimers
FORWARD: TAA TAC GAC TCA CTA TAG GG
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/clone lib="CHORI-230 Segment 1"
/clone lib="CHORI-230 Segment 1"
/note="Vector: pTARBAC2.1; Site_1: EcoRI; Site_2: EcoRI;
CHORI-230 Rat (BN/SsNHsd/MCW) BAC library produced by
Pieter de Jong"
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/mol_type="genomic DNA"
/setrain="BN/SsMHsd/MCW"
/db_xref="taxon:10116"
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90.9%;
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University of Arizona
Forbes Building Room 303, Tu
Tel: 520 626 9595
Fax: 520 621 1259
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Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
1 (bases 1 to 731)
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731 bp DNA linear GSS 26-JUL-2004
OR BBa0048E12.f OR BBa Oryza rufipogon genomic clone OR BBa0048E12
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BACKWARD: CAC TCA TTA GGC ACC CCA
INSERT Length: 161 Std Error: 0.00
Plate: 0048 row: E column: 12
Seg primer: TAA TAC GAC TCA CTA TAG GG
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Unpublished (2004)
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Kudrna, D., Muller, C., Hatfield, J.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Rod A. Wing
                                                                                                                                                                                                                                                       Class:
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78.3%; ilarity 90.9%; Conservative
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/tlssue_type="young leaves"
/lab_host="DH10B-T1 phage resistant"
/clone_llb="OR_BBa"
/note="Vector: pAGIBAC1; Site_1: HindIII; Site_2: HindIII"
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/mol_type="genomic DNA"
/db_xref="taxon:4529"
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                                                                                /tissue_type="young leaves"
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/lab_nost="OR_BBa"
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                                                                                                                                                                       /mol_type="genomic DN
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                                                                                                                                                                                                     organism="Oryza rufipogon"
                                                                                                                                                                                                                                   ocation/Qualifiers
                                                                                                                                                     clone="OR_BBa0048E12"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GI:50611301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   78.3%;
90.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                303, Tucson, AZ 85721-0036,
Score 18.8; DB 9;
Pred. No. 3.5e+02;
0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 18.8; DB 9;
Pred. No. 3.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24
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                                                                                                                                                                                        DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rao, K., Luo, M., Jetty, R., Soderlund, C. and Wing, R
                                 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 632;
                                 Length 731;
                                                                                       HindIII;
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Indels

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Gaps

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AUTHORS
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CL807004/c
LOCUS
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KEYWORDS
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                          ORGANISM
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Best Local
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                                                                                                                                                                                                                                                        152 TATGTCGACTCGTCTTCAAAG 131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       151
                                                          ;, mkNA seq
AI332619
AI332619.1
EST.
                                                                                                       926 bp mRNA linear EST 13-FEB-1999 qq28a04.x1 Soares NhHMPu S1 Homo sapiens cDNA clone IMAGE:1933806 3' similar to SW:MDP1_RABIT P31429 MICROSOMAL DIPEPTIDASE PRECURSOR 1, mRNA sequence.
                      Homo sapiens
                                                                                                                                                                                                                                                                           3 TATGTAGACACGTCTTTCAAAG 24
                                                                                                                                                                                                                                                                                                                                20;
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                       Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: http://genome.arizona.edu
PCR PRimers
FORWARD: TAA TAC GAC TCA CTA TAG GG
BACKWARD: CAC TCA TTA GGC ACC CCA
Plate: 0020 row: P column: 06
Seq primer: TAA TAC GAC TCA CTA TAG GG
Class: BAC ends.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Arizona Genomics Institute
University of Arizona
Porbes Building Room 303, Tucson,
Tel: 520 626 9595
Fax: 520 621 1259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oryza rufipogon
Oryza rufipogon
Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
1 (bases 1 to 765)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CL807004 765 bp DNA linear COR CB80020P06.f OR CB8 Oryza rufipogon genomic clone 5', genomic survey sequence.
CL807004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Rod A. Wing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CL807004.1 GI:51044586 GSS.
                                                                                                                                                                                                                                                                                                                                               Similarity
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                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                 /clone="or_CBa0020F06"
/tlssue_type="young leaves"
/dev_stage="2 week old seedlings"
/lab_host="DH10B T1 phage resistant"
/clone_1ib="OR_CBa"
/note="Vector: pAGIBAC1; Site_1: HindIII; Site_2: HindIII;
drk treated 36 hrs before harvest"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mol_type="genomic DN/
/db_xref="taxon:4529"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Oryza rufipogon"
/mol_type="genomic DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ocation/Qualifiers
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                                                                           GI:4069178
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                                                                                                                                                                                                                                                                                                                                               78.3%;
                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                          Score 18.8; DB 9;
Pred. No. 3.5e+02;
0; Mismatches 2
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                                                                                                                                                                                                                                                                                                                                                          Length 765;
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e OR_CBa0020P06
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JOURNAL
COMMENT
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BH812962/c
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ORGANISM
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DEFINITION
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AUTHORS
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Best Local S
Matches 19
                                                                                                                                                                                                                                               TITLE
                                                                                                                                                                                                                                                                                                                      AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  474
                                                                                                                                                                                                                                     1 (bases 1 to 180)
Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R.,
Gadrinab,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L.,
Shinn,P., Zimmerman,J. and Ecker,J.R.
Shopence-Indexed Library of Insertion Mutations in the
                                                                                                                    Unpublished (2001)
Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory
The Salk Institute for Biological Studies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  180 bp DNA linear GSS 02-NSALK_063510 Arabidopsis thaliana TDNA insertion lines Arabid thaliana genomic clone SALK_063510, genomic survey sequence BH812962
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             w
                                               10010 N. Torrey Pines Road,
Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                     Arabidopsis thaliana (thale cress)
Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GSS
                                                                                                                                                                                                                      Arabidopsis Genome
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Contact: Robert Strausberg, Ph.D.
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov .
This clone is available royalty-free through LLNL; contact the This Consortium (info@image.llnl.gov) for further information.
IMAGE Consortium (info@image.llnl.gov) for further information.
IMAGE Theographic Il31 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            High quality sequence stop: 412.
Location/Qualifiers
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
Tumor Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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1 (bases 1 to 526)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TATGTGGACACGTCTTTCAA 493
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  is single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone_lib="Soares NhHMPu_S1"
/note="Organ: mixed (see below); Vector: pT7T3D-Pac
/note="Organ: mixed (see below); Vector: pT7T3D-Pac
(Pharmacia) with a modified polylinker; Site_1: Not I;
Site_2: Eco RI; Equal amounts of plasmid DNA from three
normalized libraries (melanocyte 2NbHM, pregnant uterus
NbHPU, and fetal heart NbHH19W) were mixed, and ss circles
were made in vitro. Following HAP purification, this DNA
was used as tracer in a subtractive hybridization
reaction. The driver was PCR-amplified cDNAs from pools of
5,000 clones made from the same 3 libraries. The pools
consisted of I.M.A.G.E. clones 260232-265223,
340488-345479, and 484488-489479."
                              ecker@salk.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /tissue_type="Pooled human melanocyte,
pregnant uterus"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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  pass
  sequence
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Pred. No. 5.3e+02;
0; Mismatches 1;
                                                                                             La Jolla,
     recovered
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from
                                                                                             92037, USA
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the left border
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lines Arabidopsis
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AUTHORS
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BZ290962/c
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Class: TDN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.

1 (bases 1 to 197)
Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R., Gadrinab,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L., Shinn,P., Zimmerman,J. and Ecker,J.R., A Sequence-Indexed Library of Insertion Mutations in the Arabidopsis Genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N
                                                                                                                                                                                                                                                                                                              At3g26850.
Class: TDN
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                                                                                                                                                                                                                                                                                                                                                         This is single pass sequence recovered from the left border of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BZ290962.1 GI:24335370
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                                                                                                                                                                                                                                                                                                                                                                                                             ecker@salk.edu
                                                                                                                                                                                                                                                                                                              TDNA tagged
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each of which contains one or more TDNA insertion
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the site of insertion. Details of the protocols used can
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elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
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ecotype="Col-0"
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87.0%;
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Pred. No. 5.8e+02;
0; Mismatches 3;
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                                                   http://signal.salk.edu/tdna_protocols.html"
 Score 18.2; DB 8;
Pred. No. 5.9e+02;
               8;
               Length 197;
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ORIGIN

片 8 Query Match
Best Local Similarity
Matches 20; Conserv 86 TTATGTAGACACGTCTTTCAAAG 24 TTATGTAGACAAGTATGTCAAAG 76 Conservative 0; Mismatches Indels 0; Gape

0

Search completed: February Job time: 2354.9 secs

9, 2005, 21:55:30

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OM nucleic - nucleic search, using sw model
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Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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GenEmbl:
1: gb ba:
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3: gb htg:
4: gb ow:
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7: gb pat:
7: gb pat:
10: gb ro:
11: gb at:
11: gb at:
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Gapop 10.0 , Gapext 1.0
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25
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Copyright (c) 1993 - 2005 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

c 19	c 18	c 17	c 16	c 15	C 14	c 13	c 12	c 11	c 10	ი •	ი 8	c 7	ი ი	n u	Ω 4	ი u	ი 2	1	Result No.
25	25	25	25	25	25	25	25	25	25	25	25	25	25	25	25	25	25	25	Score
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AX598697	AX457064	CQ806551	HUMMDR1A	A38669	AR080192	AR080193	HUMMDR1P	AR080194	G19985	HUMMDR1A02	AX701741	AR080195	AF345624	AF345623	AR080197	AR080196	AX504324	AX504307	ID
AX598697 Sequence	AX457064 Sequence	CQ806551 Sequence	M57450 Human multi	A38669 Sequence 1	AR080192 Sequence	AR080193 Sequence	L07624 Human multi	AR080194 Sequence	G19985 BWSS10 Eric	M29423 Human P-gly	AX701741 Sequence	AR080195 Sequence	AF345624 Homo sapi	AF345623 Homo sapi	_	AR080196 Sequence	AX504324 Sequence	AX504307 Sequence	Description

ALIGNMENTS

RESULT 2 AX504324/c LOCUS DEFINITION ACCESSION VERSION KEYWORDS SOURCE ORGANISM	Qy Db	Query Match Best Local : Matches 2	Source ORIGIN	AUTHORS TITLE JOURNAL	SOURCE SOURCE ORGANISM	RESULT 1 AX504307 LOCUS DEFINITION ACCESSION VERSION
AX504324 25 bp DNA linear PAT 27-SEP-2002 Sequence 27 from Patent WO0234291. AX504324 AX504324.1 GI:23386136	1 GTGCTCAGCCCACGCCCCGGCGCTG 25	100.0%; Score 25; DB 6; Length 25; Similarity 100.0%; Pred. No. 41; 5; Conservative 0; Mismatches 0; Indels 0; Gaps		Colgan, S.P. Compositions and methods for treating hematologic malignancies and multiple drug resistance multiple drug resistance Patent: WO 0234291-A 10 02-MAY-2002; THE BRIGHAM AND WOMEN'S HOSPITAL, INC. (US)	Homo sapiens (human) Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.	AX504307 25 bp DNA linear PAT 27-SEP-2002 Sequence 10 from Patent WO0234291. AX504307 AX504307.1 GI:23386125
N		0;				8

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                                                          1 (bases 1 to 351)
Stein, U. and Walther, W.
Vector for the expression of therapy-relevant
Patent: US 5968735-A 6 19-OCT-1999;
                                                                                                                      Unknown.
                                                                                                                                    Unknown.
                                                                                                                                                                        Sequence 6 from patent US AR080197
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Stein, U. and Walther, W.
Vector for the expression of therapy-relevant genes
Patent: US 5968735-A 5 19-OCT-1999;
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AR080196
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Patent: WO 0234291-A 27 02-MAY-2002;
THE BRIGHAM AND WOMEN'S HOSPITAL, INC.
Location/Qualifiers
                                                                                                            Unclassified.
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Mammalia; Eutheria;
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|larity 100.0%; !
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                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 370)

Alberta, E.O., Petersen, R.H., Hughes, K.W. and Lechner, B. Miscellaneous notes on Pleurotus
Persoonia 18, 55-69 (2002)

2 (bases 1 to 370)
                                                                                                                             Homo sapiens
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AF345624.1
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Homo sapiens MES-SA/VL20-4.2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (06-FBB-2001) Oncology/Medicine, Stanford University
School of Medicine, 269 Campus Drive, Stanford, CA 94305-5151, USA
Location/Qualifiers
Wang,Y., Chen,G.K. and Sikic,B.I. Direct Submission
                                                                                                                                         Homo sapiens (human)
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Submitted

(06-FEB-2001) Oncology/Medicine, Stanford University

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Sequence 3 from Patent W003002760.
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Location/Qualifiers
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                             Distler, J. and Leu, E. Method for detecting cytosine methylation analysing single strands of amplificates patent: WO 03002760-A 3 09-JAN-2003;
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/mol_type="mRNA"
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/chromosome="7"
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/note="vinblastine selected cell line"
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/mol_type="unassigned
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M29423 J05168 M18754
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90094448
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J. Biol. Chem. 265 (1), 506-514 (1990)
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Isolation and sequence of the promoter region
multidrug-resistance (P-glycoprotein) gene
J. Biol. Chem. 262 (36), 17432-17436 (1987)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Chen, C.J., Clark, D., Ueda, K., Pastan, I., Gottesman, M.M.
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/db_xref="taxon:9606"
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Unpublished (1997)
Synonyms: PGY1
GDB: GDB:583426
GDB_DSEG: PGY1
Contact: Eric D. Green
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                                                                                                                      This STS was developed from sequence determined by anot investigator. See GenBank record: M29423 For additional information about the NHGRI chromosome 7 mapping project, http://www.nhgri.nih.gov/DIR/GTB/CHR7. Also see Genomics 11:548-64 (1991) [MUID=92128937]
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National Human Genome Research
49 Convent Dr., MSC4431, Bldg.
Tel: 3014020201
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Bouffard,G.G., Iyer,L.M., Idol,J.R., Braden,V.V., Cunr
Bouffard,G.A., Mohr-Tidwell,R.M., Peluso,D.C., Fultor
Weintraub,L.A., Mohr-Tidwell,R.M., Peluso,D.C., Fultor
Leckie,M.P. and Green,E.D.
A collection of 1814 human chromosome 7-specific STSs
Genome Res. 7 (1), 59-64 (1997)
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sWSS10 Eric D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: egreen@nhgri.nih.gov
Primer A: AACGGAAGCCAGAACATTCC
Primer B: AGGCTTCCTGTGGCAAAGAG
STS size: 180
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                /organism="Homo sapiens"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                976 bp DNA linear STS 28-SEP sapiens STS genomic, sequence tagged
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Institute/NIH
49, Rm. 2A08, Bethesda, MD 20892
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                                                                                                                                                                                                                                                                                                                                                                                                  T.
                                                                                                                                                                                                                                                                                                                                                                                                                              for
                                                                                                                                         mapping project, see Also see Genomics
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2.00
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minute(s)
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minute(s)
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HUMMDR1P/c
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AUTHORS
TITLE
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ORGANISM
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AR080194/c
LOCUS
DEFINITION
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Matches
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                                                                                                                                        JOURNAL
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Best Local Similarity
Matches 25; Conserv
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1 (bases 1 to 1318)

1 (bases 1 to 1318)

Stein, U. and Walther, W.

Vector for the expression of therapy-relevant genes

Patent: US 5968735-A 3 19-OCT-1999;

Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 3
AR080194
AR080194.1
                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 1327)

Madden,M.J., Morrow,C.S., Nakagawa,M., Goldsmith,M.E.,

Pairchild,C.R. and Cowan,K.H.

Identification of 5' and 3' sequences involved in the regulation of transcription of the human mdrl gene in vivo

3. Biol. Chem. 268 (11), 8290-8297 (1993)
                                                                                                                                                                                                                                                     L07624.1 GI:187500
MDR1 gene; P-glycoprotein; multidrug resistance protein.
Homo sapiens (human)
Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25;
                                                                           Original source text: Homo sapiens 4a) bone marrow DNA.
                                                                                                           8096520
                                                                                                                                                                                                                                                                                                                                                Human multidrug resistant P glycoprotein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
/organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
                                              Location/Qualifiers
1. .1327
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/gene="PGY1"
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253. .432
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="unknown"
/mol_type="unassigned
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from patent US 5968735.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 25; DB
100.0%; Pred. No. 16;
tive 0; Mismatches
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Pred. No. 17;
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PGY1)

PRI 09-JAN-1995 L) gene, 5'

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DEFINITION
ACCESSION
VERSION
KEYWORDS
SOURCE
ORGANISM
                                                REFERENCE
AUTHORS
TITLE
JOURNAL
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AR080193/c
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                                                                                                                                           1688 bp
Sequence 2 from patent US 5968735.
AR080193
AR080193.1 GI:1000692A
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                               1 (bases 1 to 1688)
Stein, U. and Walther, W.
Vector for the expression of therapy-relevant genes
Patent: US 5968735-A 2 19-OCT-1999;
Location/Qualifiers
                                                                                                              Unknown.
                                                                                                                          Unknown
                                                                                                Unclassified.
                                                                                                                                                                                                                                                                                                                 Similarity
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llarity 100.0%;
Conservative 0
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/tissue type="bone marrow"
/tissue_lib="lambda Charon
863. .1171
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/note="G00-120-712"
/bound_moiety="unknown"
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/bound moiety="unknown"
function="heat shock element"
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/bound_moiety="unknown"
/function="heat_shock_element"
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/gene="PGY1"
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|041. .1170
|gene="PGY1"
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.041. .>1170
'gene="PGY1"
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/mol_type="unassigned DNA"
          organism="unknown"
                                                                                                                                                                                                                                                                                                                                                                              note="G00-120-712"
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                          .1688
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Pred. No. 16;
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RESULT 14
AR080192/c
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DEFINITION
ACCESSION
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AUTHORS
TITLE
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Matches 25
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Best Local S
Matches 25
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TITLE
JOURNAL
Query Match
Best Local Similarity
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                                                                                      variation
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AR080192
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Location/Qualifiers
                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primatee; Catarrhini; Hominidae; Homo.

1 (bases 1 to 2117)
Stein, U. and Walther, W.
VECTOR FOR THE EXPRESSION OF THERAPY-RELEVANT GENES
PATENT: WO 9411522-A 1 26-MAX-1994;
MAX DELBRUECK CT PUER MOLEKULA (DE)
Other publication DE 4238778 940519.
Location/Qualifiers
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A38669
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ilarity 100.0%;
Conservative 0
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ilarity 100.0%; Pred. No. 15
Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="unknown"
/mol_type="unassigned
                                                /replace="t"
                                                                                                                                     organism="Homo sapiens"
/mol_type="unassigned DN/db_xref="taxon:9606"
                                                                        replace="c"
                                                                                                 cell_type="SARCOMAS"
tissue_type="TUMORS"
                                                                                                                            clone="CVS-SW1"
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100.0%;
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patent US 5968735.
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Pred. No.
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W09411522.
  Score 25;
Pred. No.
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  DB
15;
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2005, 17:01:54	25; Conservative 0; Mismatches 1 GTGCTCAGCCCAGGCCCCGGCCCTG 25
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	Indels
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Result
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Maximum DB
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score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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1: /cgn2_6/ptcdata/1/ina/5A_COMB.seq:*

2: /cgn2_6/ptcdata/1/ina/5B_COMB.seq:*

3: /cgn2_6/ptcdata/1/ina/6A_COMB.seq:*

4: /cgn2_6/ptcdata/1/ina/6B_COMB.seq:*

5: /cgn2_6/ptcdata/1/ina/backfiles1.seq:*
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Copyright (c) 1993 - 2005 Compugen Ltd.
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US-08-439-814-3
US-08-439-814-3
US-08-439-814-3
US-08-439-814-2
US-08-439-814-1
US-08-181-471-2
US-09-023-655-1167
US-08-58-3-276-18
5206352-3
US-08-52-447-1
US-09-316-167-1
US-09-316-167-1
US-09-927-561-2
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US-09-925-40-1266
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Sequence 11941, Ap
Sequence 174342,
Sequence 117434,
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Sequence 19, Appl	Sequence 19, Appl	, L	, ,	, 17	17,	sequence I/, Appr	17	11,	Sequence I, Appli	ما ه	Sequence 3, Appli	, -	GENERAL INFORMA	٠,	· -	-	Sequence 1, Appli

ALIGNMENTS

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US-08-439-814-5/c
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CURRENT APPLICATION NUMBER: US/08/439,814
FILING DATE: 12-MAY-1995
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: DE P4238778.7
FILING DATE: 12-NOV-1992
PRIOR APPLICATION NUMBER: DE PCT/DE93/01086
FILING DATE: 10-NOV-1993
ATTORNEY/AGENT INFORMATION:
NAME: KLESNER, Sharon N.
REGISTRATION NUMBER: 91614-5015
REFERENCE/DOCKET NUMBER: P1614-5015
REJECOMMUNICATION INFORMATION:
TELEPHONE: 202/638-5000
TELEFAX: 202/638-6000
TELEFAX: 202/638-6000
TELEFAX: P1609Y disk
COMPUTER READABLE FORM:
MEDIUM TYPE: F10py disk
COMPUTER: IBM PC COMPATION
PRIOR APPLICATION NUMBER: DC-DOS/MS-DOS
SOFTWARRE: Patentin Release #1.0, Version
PRIOR APPLICATION NUMBER: DE P4238778.7
FILING DATE: 12-NOV-1992
PRIOR APPLICATION NUMBER: WO PCT/DE93/01086
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APPLICANT: STEIN, Ulrike
APPLICANT: WALTHER, Wolfgang
TITLE OF INVENTION: VECTOR FOR THE EXPRESSION
TITLE OF INVENTION: THERAPY-RELEVANT GENES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 20005-5701
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
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W., Suite 330
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RESULT 2
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                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/439,814
FILING DATE: 12-MAY-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE P. 4238778.7
FILING DATE: 12-NOV-1992
PRIOR APPLICATION NUMBER: DE PCT/DE93/01086
FILING DATE: 10-NOV-1993
ATTORNEY/AGENT INFORMATION:
NAME: KLESNER, Sharon N.
REGISTRATION NUMBER: 36,335
REGERENCE/DOCKET NUMBER: P1614-5015
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202/638-5000
TELEPAX: 202/638-4810
TELEPAX: 202/638-4810
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, V6
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
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APPLICANT: STEIN, Ulrike
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NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 351 base pairs
                                                                  PRIOR APPLICATION UNBER: WO PCT/DE93/01086
                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: STEIN, Ulrike
APPLICANT: WALTHER, Wolfgang
TITLE OF INVENTION: VECTOR FOR THE EXPRESSION OF
TITLE OF INVENTION: THERAPY-RELEVANT GENES
                                                                                                                                APPLICATION DATA:
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                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30B (EPO)
                                                      APPLICATION NUMBER: WO PFILING DATE: 10-NOV-1993
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Local Similarity 100.0%; Pred. No. 0.9
es 25; Conservative 0; Mismatches
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Pred. No. 0.93;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; MOLECULE TYPE: DNA (genomic) US-08-439-814-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 4, Applic Patent No. 5968735
                                                                                           FILING DATE: 10-NOV-19
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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FILING DATE: 12-NOV-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE PCT
APPLICATION ONDER: DE PCT
FILING DATE: 10-NOV-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                       APPLICATION NUMBER: WO PCT/DE93/01086
                                                                                                                                                                                                         SOFTWARE: PatentIn Release #1.0, 1
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE P4238778.7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 20005-5701
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: (FILING DATE: 12-MAY-: CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTWARB: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: STEIN, Ulrike
APPLICANT: WALTHER, WOLFGANG
TITLE OF INVENTION: VECTOR FOR THE EXPRESSION
TITLE OF INVENTION: THERAPY-RELEVANT GENES
   MOLECULE TYPE:
                                                                       LENGTH:
                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION: TELEPHONE: 202/638-5000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                      NAME: KLESNER, Sharon N. REGISTRATION NUMBER: 36,335 REFERENCE/DOCKET NUMBER: P1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 655 Fifteenth Street, N. W., Suite 330 G STREET: Street Lobby CITY: Washington STATE: DC
                                                   TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                              TELEPHONE: 202/000
TELEPHONE: 202/638-4810
                                 STRANDEDNESS: single
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mes 25; Conservative
                     COPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: singl
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DNA (genomic)
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                                                            US-08-439-814-3
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Best Local
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                                                                                                                                                 PILING DATE: 12-NOV-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PC'
PILING DATE: 10-NOV-1993
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 1318 base pairs
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/439,814
FILING DATE: 12-MAY-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE P 4238778.7
PRILING DATE: 12-NOV-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE PCT/DE93/01086
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 10-NOV-1993
ATTORNEY/AGENT INFORMATION:
NAME: KLESNER, Sharon N.
REGISTRATION NUMBER: 36,335
REFERENCE/DOCKET NUMBER: P1614-5015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS: NIKAIDO, MARMELSTEIN, MURRAY & ORAM LLP
ADDRESSES: NIKAIDO, MARMELSTEIN, MURRAY & ORAM LLP
STREET: 655 Fifteenth Street, N. W., Suite 330 G
STREET: Street Lobby
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                              TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0, Version #1.30B (EPO)
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: 202/638-500
TELEFAX: 202/638-4810
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy di
                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION: TELEPHONE: 202/638-5000
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TITLE OF INVENTION: THERAPY-RELEVANT GENES
TITLE OF INVENTION: THERAPY-RELEVANT GENES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
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CITY: W
                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                   STRANDEDNESS: single
                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                414 GrecreAGCCCACGCCCGGCGCTG 390
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                                                                                                     linear
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                                                                            DNA (genomic)
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100.0%; Pred. No. 0.91;
    100.0%;
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    Score 25; DB 2; Pred. No. 0.87;
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                      Length 1318;
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                                                                ; MOLECULE TYPE: US-08-439-814-2
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Patent No. 5968735
GENERAL INFORMATION:
APPLICANT: STEIN, Ulrike
APPLICANT: WALTHER, Wolfgang
TITLE OF INVENTION: VECTOR FOR THE EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
Best Local
Matches
                                Query Match
                                                                                                                                                                     FILING DATE: 10-NOV-1993
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: DE P 4238778.7
FILING DATE: 12-NOV-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE PCT/DE93/010
FILING DATE: 10-NOV-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 202/638-4810
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30B (EPO)
PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: NIKAIDO, MARMELSTEIN, MURRAY & ORAM LLP STREET: 655 Fifteenth Street, N. W., Suite 330 G STREET: Street Lobby
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 20005-5701
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FILING DATE: 12-NOV-1992
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REFERENCE/DOCKET NUMBER: P1614-5015
                                                                                                        TOPOLOGY:
                                                                                                                    STRANDEDNESS
                                                                                                                                   LENGTH: 1688 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                COMPUTER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: KLESNER, Sharon N
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                                                                                                                                                                                                                           APPLICATION NUMBER: WO PCT/DE93/01086
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                   Similarity
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 100.0%; ilarity 100.0%; Conservative 0
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                                                                                    DNA (genomic)
                                                                                                                      single
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                   Score 25; DB 2
Pred. No. 0.86;
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     Mismatches
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                                       DB 2;
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                                   Length 1688;
       Indels
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US-08-439-814-1/c

Sequence 1, Applic Patent No. 5968735

Application US/08439814

GENERAL INFORMATION:

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1534 GTGCTCAGCCCACGCCCCGGCGCTG 1510

US-08-181-471-2/c

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RESULT 7
                                                                                                                               Matches
                                                                                                                                                                 Query Match
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ZIP: 20005-5701
COMPUTER READABLE FORM:
COMPUTER TEADABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: IBM PC compatible
TYPHTER: IBM PC compatible
TYPHTER: TOTAL PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                        FILING DATE: 10-NOV-1993
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:

APPLICATION NUMBER: DE P 4
FILING DATE: 12-NOV-1992
PRIOR APPLICATION DATA:
APPLICATION UMBER: DE PCT
APPLICATION UMBER: DE PCT
FILING DATE: 10-NOV-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                         MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Patentin Release #1.0, 1
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE P4238778.7
PILING DATE: 12-NOV-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: STELN, WALTHER, WOLIGANG
APPLICANT: WALTHER, WOLIGANG
TITLE OF INVENTION: THERAPY-RELEVANT GENES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 12-MAY-19
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: NIFTCH OF MARMELSTEIN, MURRAY & ORAM LLP STREET: 655 Fiftcenth Street, N. W., Suite 330 G STREET: Street Lobby
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: 202/638-5000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET:
STREET:
CITY: W
                                                                                                                                              Local
                                                                                                                                                                                                                                                       STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: 202/638-50
TELEFAX: 202/638-4810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: KLESNER, Sharon N.
REGISTRATION NUMBER: 36,335
REFERENCE/DOCKET NUMBER: P1614-5015
                                                                                                                                                                                                                                            : YOULOGY:
                                                                                                                                                                                                                                                                                             ENGTH:
                                                       1936 GTGCTCAGCCCACGCCCCGGCGCTG 1912
                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                          1: 2090 base pairs nucleic acid
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                                                                                         GTGCTCAGCCCACGCCCCGGCGCTG 25
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                                                                                                                                                                                                                     DNA (genomic)
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                                                                                                                                                                                                                                                     single
                                                                                                                           100.0%; Score 25; DI
100.0%; Pred. No. 0.0
Live 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Version #1.30B (EPO)
                                                                                                                                              0.85;
                                                                                                                                                              DB 2;
                                                                                                                           0;
                                                                                                                                                              Length 2090;
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US-09-023-655-1167/c
; Sequence 1167, Application US/09023655
; Patent No. 6607879
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                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 619-792-8477
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
      CORRESPONDENCE ADDRESS:
ADDRESSEE: INCYTE PHARMACEUTICALS, INC.
STREET: 3174 PORTER DRIVE
                                                                                                       APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/181,471

FILING DATE: 13-JAN-1994
                                                                       TITLE OF INVENTION:
                                                          NUMBER OF SEQUENCES:
                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                             ANTI - SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Li, Lin
APPLICANT: Lishko,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 02-APR-1993
TORNEY/AGENT INFORMATION:
NAME: Fitting, Thomas
REGISTRATION NUMBER: 34,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: AN LECOMMUNICATION INFORMATION: TELEPHONE: 619-792-3680
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: San Diego
                                                                                                                                                                                                                                                                                                                                                                                                  LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Thomas Fitting
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                                                                                                                                                                                                                                                           251 GTGCTCAGCCCACGCCCCGGCGCTG 227
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PALO ALTO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nucleic acid
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                                                                                                     Susan G. Stuart
Jeffrey J. Seil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4646 base pairs
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425..4267
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                                                                       J. Seilhamer
COMPOSITION
EXPRESSION
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METHOD FOR DELIVERING BENEFICIAL
METHOD FOR DELIVERING BENEFICIAL
COMPOSITIONS TO HAIR FOLLICLES
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Pred. No. 0.82;
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                                                                                         FOR THE DETECTION OF BLOOD CELL GENE
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CALIFORNIA

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US-08-583-276-18/c
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US-09-023-655-1167
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Zeller, Karen J.
REGISTRATION NUMBER: 37,071
REFERENCE/DOCKET NUMBER: PA-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (650) 845-416
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
              COMPUTER: IBM PS/2
ODERATING SYSTEM: PC-DOS
SOFTWARE: DW4.V2
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/
FILING DATE: 05-JAN-1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS 6.2
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/023,655
FILING DATE: HEREWITH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IMMEDIATE SOURCE:
LIBRARY: GENBA
                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch diskette
                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: McDonagh, Kevin T.
APPLICANT: Nienhuis, Arthur
APPLICANT: Tolstoshev, Paul
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: CALIFO
COUNTRY: USA
                                                                                                                                                                                                                                                            ADDRESSEE: Carella, Byrne, Bain, Gilfillan
ADDRESSEE: Cecchi & Stewart
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                                                                                                             CITY: Roseland
                                                                                                                                                                                                                                            STREET: 6 Becker Farm Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS:
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                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Match 100.0%; Score 25; DB 4; Length 4646; Local Similarity 100.0%; Pred. No. 0.82;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ENGTH:
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5837536
                                                                                                                                                                                                            New Jersey
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                                                                                                                                                                                                                                                                                                                                    IMPROVED EXPRESSION OF HUMAN
MULTIDRUG RESISTANCE GENES AND IMPROVED
SELECTION OF CELLS TRANSDUCED WITH SUCH GENES
                                                                                                                                                                                                                                                                                                                      19
                                                   US/08/583,276
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RESULT 11
5206352-3/c
;Patent No.
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5206352-3/c
;Patent No. 5206352
; AppLICANT: Roninson, Igor B.;Pastan Ira H.;Gottesman,
; Michael M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               밁
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                                  5206352-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: COMPOSITIONS FOR CLONES CONTAINING DNA SEQUENCES ASSOCIATED WITH MULTIDRUG RESISTANCE IN HUMAN CELLS
                                                                                                                                                                                                                                                      Michael M.
TITLE OF INVENTION: COMPOSITIONS FOR CLONES CONTAINING DNA; SEQUENCES ASSOCIATED WITH MULTIDRUG RESISTANCE IN HUMAN CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO:3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 08, FILING DATE: 31-0CT-19; APPLICATION NUMBER: 07, FILING DATE: 22-MAY-19; INFORMATION FOR SEQ ID NO:
                                                                    SEQ ID NO:3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/
FILING DATE: 24-SEP-1990
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE:
DESCRIPTION: Genomic DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 4669 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 4
CURRENT APPLICATION DATA:
                                                                                                                                                            APPLICATION NUMBER: US/
FILING DATE: 24-SEP-1990
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                           APPLICANT: Roninson, Igor B.; Pastan Ira H.; Gottesman,
                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: singular
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 845,610 FILING DATE: 28-MAR-1986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 4669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 892,575 FILING DATE: 01-AUG-1986
                                                                                                          APPLICATION NUMBER: 845,610
                                                                                                                          APPLICATION NUMBER: 892,575 FILING DATE: 01-AUG-1986
                                                     LENGTH: 4669
                                                                                     FILING DATE: 28-MAR-1986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ocal Similarity
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22-MAY-1992
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Length 4669;
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Query Match

100.0%;

Score 25;

B 6

Best Local Similarity

100.0%;

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RESULT 13
US-09-316-167-1/c
                                                                                                  吊
                                                                                                                                                                                                               NAME/KEY:
LOCATION:
US-08-752-447-1
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US-08-752-447-1/c
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                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 312-913-9808
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 4669 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
           Sequence 1, Application US/09316167
Patent No. 6365357
                                                                                                                                                        Matches
                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA
APPLICATION UMMBER: US/08/752,447
FILING DATE: 15-NOV-1996
CLASSIFICATION: 435
ATTORNEY AGENT INFORMATION:
NAME: NO. 599408BTAID, Kevin E
REGISTRATION NUMBER: 35,303
REFERENCE/DOCKET NUMBER: 35,121
TELECOMUNICATION INFORMATION:
TELEPHONE: 312-913-0001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1, Applic
Patent No. 5994088
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
GENERAL INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: FC-DOS/MS-DOS
                                                                                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADUAL STREET: 300
CITY: Chicago
CITY: Illinois
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                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                NAME/KEY:
LOCATION:
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                                                                                               251 GCGCTCAGCCCACGCCCGGCGCTG 227
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                                                                                                                   GTGCTCAGCCCACGCCCCGGCGCTG 25
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                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                      CDS
425..4264
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1..424
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0 South Wacker Drive, Seventh Floor
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                                                                                                                                                     Score 23.4; D. Pred. No. 3.3; 0; Mismatches
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                                                                                                                                                                                   2; Length 4669;
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for P-glycoprotein
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PILING MALE:
CLASSIFICATION:
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/752,447
FILING DATE: 15-NOV-1996
ATTORNEY/AGENT INFORMATION:
NAME: No. 6365357nan, Kevin E
REGISTRATION NUMBER: 35,303
REFERENCE/DOCKET NUMBER: 95,11
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-913-0001
TELEFAX: 312-913-908
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 4669 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                       US-09-397-233-1/c
; Sequence 1, Application US/09397233
; Patent No. 6630327
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                  RESULT 14
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Matches
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STREET:
CITY: Chicago
CITY: Illinois
TISA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                     Local
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LOCATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OCATION:
                                                                                  TITLE OF INVENTION: Methods and Reagents for Preparing and Title OF INVENTION: Methods and Reagents Specific for P-glycoprotein
CORRESPONDENCE ADDRESS:
ADDRESSEE: McDonnell Boehnen Hulbert
STREET: 300 South Wacker Drive
CITY: Chicago
STATE: Illinois
                                                                                                                                                       APPLICANT: Mechetner, Eugene
                                                                                                                                                                                                                                                                                                   251
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300 South Wacker Drive, Seventh Floor
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TENTION: Methods and Reagents for Preparing
TENTION: Using Immunoligcal Agents Specific
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COUNTRY: USA
ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk.

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US-08-487-1418-2
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Matches
                                                                                                                                                                                                                                                                                                                                                                            Sequence 2, Application US/08487141B Patent No. 5683987
GENERAL INFORMATION:
APPLICANT: Smith, Larry J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                 TITLE OF INVENTION: Therapeuti
TITLE OF INVENTION: Targeting
NUMBER OF SEQUENCES: 114
CORRESPONDENCE ADDRESS:
                              SOPTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/487,141B FILING DATE: 07-UN-1995 CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATION GATA:
OPERATION DATA:
SOCIETY
OPERATION NUMBER: US/09/397,233
FILING DATE: 16-Sep-1999
CLASSIFICATION: UNMBER: US/09/397,233
FILING DATE: 16-Sep-1999
CLASSIFICATION: UNMBER: US/09/397,233
FILING DATE: NFORMATION:
NAME: No. 6630327nan, Kevin E
REGISTRATION NUMBER: 35,303
REFERENCE/DOCKET NUMBER: 95,1121-C
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-913-0001
TELEPAX: 312-913-0002
                                                                                                                                                                                                                 STREET: 1601 Market Street Suite 720
CITY: Phadelphia
STATE: PA
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1es 24; Conservative
APPLICATION NUMBER:
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LOCATION: 4265..4669
SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              251 GCGCTCAGCCCACGCCCGGCGCTG 227
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 4669 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LOCATION:
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425..4264
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1..424
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                                                                                                                                                                                                                                                                                                                                            Therapeutic Oligonucleotides
Targeting the Human MDR1 and
US 08/379,180
                                                                                                                                                                                                                                                                                                                                              MRP Genes
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; ANTI-SENSE: US-08-487-1418-2
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                                                                 Query Match
Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                  TELEPHONE: (215)563-4100
TELEPAX: (215)563-4044
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                   FILING DATE: 12-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Hagan, Patrick J.
REGISTRATION NUMBER: 27,643
REFERENCE/DOCKET NUMBER: 6308
TELECOMMUNICATION INFORMATION:
                                                                                                                                                     MOLECULE TYPE: DN
HYPOTHETICAL: NO
ANTI-SENSE: YES
                                                                                                                                                                                                    LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: not relevant
                                   1 GTGCTCAGCCCACGCCCCGG 20
GTGCTCAGCCCACGCCCCGG
                                                                    Conservative
                                                                                                                                                                                                                                                                                                                        (215) 563-4100
                                                                                                                                                                                    DNA (genomic)
                                                                 80.0%; Score 20; DB
100.0%; Pred. No. 82
ative 0; Mismatches
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 20
                                                                                    DB 1; Length 20; 82;
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Search completed: February 9, 2005, 17:11:14 Job time : 79.592 secs

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Database
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Listing first 45 summaries
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Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Title:
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                                                                                                                                                                                                                                                                                                                                                                                    Published Applications NA:*

1: /cgn2 6/ptodata/1/pubpna/PCT NEW PUB.seq:*

2: /cgn2 6/ptodata/1/pubpna/PCT NEW PUB.seq:*

3: /cgn2 6/ptodata/1/pubpna/USO6 NEW PUB.seq:*

4: /cgn2 6/ptodata/1/pubpna/USO6 PUBCOMB.seq:*

5: /cgn2 6/ptodata/1/pubpna/USO7 NEW PUB.seq:*

6: /cgn2 6/ptodata/1/pubpna/PCTUS PUBCOMB.seq:*

7: /cgn2 6/ptodata/1/pubpna/USO8 NEW PUB.seq:*

8: /cgn2 6/ptodata/1/pubpna/USO8 PUBCOMB.seq:*

9: /cgn2 6/ptodata/1/pubpna/USO8 PUBCOMB.seq:*

9: /cgn2 6/ptodata/1/pubpna/USO8 PUBCOMB.seq:*
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Gapop 10.0 , Gapext 1.0
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25
                                                                                                                                                                                                                                                                       4313806 seqs, 2877871033 residues
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Copyright (c) 1993 - 2005 Compugen Ltd
/ Cgn12 6/ptodata/1/pubpna/US09A PUBCOMB.seq: *
/ Cgn2 6/ptodata/1/pubpna/US09B PUBCOMB.seq: *
/ Cgn2 6/ptodata/1/pubpna/US09B PUBCOMB.seq: *
/ Cgn2 6/ptodata/1/pubpna/US09D PUBCOMB.seq: *
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/ Cgn2 6/ptodata/1/pubpna/US10D PUBCOMB.seq: *
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(without alignments)
539.601 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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Score 25 2 25 2 25 3 25	2!	2	2	5 2	7 2!	8 2!	9 2!	2	2!	
a Match Length DB ID 100.0 633 18 U 5 100.0 2307 9 US 5 100.0 2932 18 U									5 100.0	
Length 633 2307 2932	2932	4533	4643	4643	4643	4646	4646	4646	4646	
18 91 18	18	ω	13	14	15	11	片	17	17	
ID US-10-482-433A-3 US-09-805-020-31 US-10-473-126-37	US-10-473-126-37	US-09-805-020-30	US-10-072-621-2	US-10-097-340-1	US-10-007-926A-258	US-09-968-007A-459	US-09-968-007A-747	US-10-641-643-1167	US-10-343-657-1	
Description Sequence 3, Appli Sequence 31, Appl Sequence 37, Appl	Sequence 37, Appl	Sequence 30, Appl	Sequence 2, Appli	Sequence 1, Appli	Sequence 258, App	Sequence 459, App	Sequence 747, App	Sequence 1167, Ap	Sequence 1, Appli	

RESULT 2 US-09-805-020-31/c

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18.2																18.2					18.4	18.4	18.4	18.6	18.6	18.6	18.6	18.6	œ	19.2	19.2	23.4	25
72.8	72.0	72.8	72.8	72.8	72.8	72.8	72.8	72.8	72.8	72.8	72.8	72.8	72.8	72.8	72.8	72.8	•	٠	'n	72.8	•	73.6	73.6	74.4	•	•	74.4	74.4	74.4	9		ω	100.0
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13	17	18	18	ω	17	18	17	17	14	14	14	15	17	17	13	18	18	16	14	18	18	17	18	16	15	13	18	18	9	H	ø	18	18
1 1	TIS-10-092-900A-153	US-10-425-115-150283	US-10-494-674-85	US-09-738-626-273	US-10-302-172-845	US-10-437-963-72302	-10-425	US-10-425-114-28838	US-10-224-260-21	US-10-224-260-19	US-10-224-260-17	US-10-241-313-3	US-10-425-114-31550	US-10-027-632-164589	US-10-027-632-164589		US-10-767-701-8299	US-10-029-386-7696	-10-224-260-1	US-10-425-115-63236	-10-322-281-544	-10-282-	-10	US-10-146-733-61	US-10-154-419-66	US-10-024-623-16	US-10-767-701-5342	US-10-357-930-56850	US-09-864-761-13425	US-09-968-007A-755	US-09-880-107-3828	Ļ	US-10-775-169-198
575,		150	Sequence 85, Appl	~	Sequence 845, App	e 72302	31892,		21,	19	17	·	ι L			Sequence 6, Appli			LINFOR		544, Ap	17493		Sequence 61, Appl							3828,	Sequence 1, Appli	Sequence 198, App

ALIGNMENTS

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SEQ ID NO 3
LENGTH: 633
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Amplification Product of MdRI-Fragment
US-10-482-433A-3
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US-10-482-433A-3/c
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Publication No. US20040265814A1

Publication No. US20040265814A1

PUBLICANT: Epigenomics AG

TITLE OF INVENTION: Method for the detection of cytosine methylation by comparative

TITLE OF INVENTION: analysis of single strands of amplificates
                                                                                            Matches
                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 82.174
CURRENT APPLICATION NUMBER: US/10/482,433A
CURRENT FILING DATE: 2003-12-29
PRIOR APPLICATION NUMBER: DE 10132212.7
PRIOR FILING DATE: 2001-06-27
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin version 3.2
                                                                                       Match 100.0%; Score 25; DB 1 Local Similarity 100.0%; Pred. No. 0.24; Pred. No. 0.24; Mismatches
422
                                               1 GTGCTCAGCCCACGCCCCGGCGCTG 25
  GTGCTCAGCCCACGCCCGGCGCTG 398
                                                                                                                                         DB 18; Length 633;
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GENERAL INFORMATION:

APPLICANT: LEVINE, Zurit

TITLE OF INVENTION: SPLICE VARIANTS OF ONCOGENES

FILE REFERENCE: 2786-0168P

CURRENT APPLICATION NUMBER: US/09/805,020

CURRENT FILING DATE: 2001-03-13

NUMBER OF SEQ ID NOS: 72

SOFTWARE: PATENTIN Ver. 2.1

SEQ ID NO 31

LENGTH: 2307
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TYPE: DNA
; ORGANISM: Homo Sapiens
US-10-473-126-37
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US-10-473-126-37/c
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                                                                                                                                                                                                  Sequence 30, Application US/09805020 Publication No. US20020086384A1 GENERAL INFORMATION:
                                                                 SEQ ID NO 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 37, Application US/10473126
Publication No. US20040234973A1
GENERAL INFORMATION:
APPLICANT: Epigenomics AG
TITLE OF INVENTION: Methods and nucleic acids for the analysis of
TITLE OF INVENTION: proliferative disorders
                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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Matches
                                                                            TITLE OF INVENTION: SPLICE VARIANTS OF ONCOGENES FILE REFERENCE: 2786-0168P CURRENT APPLICATION NUMBER: US/09/805,020 CURRENT FILING DATE: 2001-03-13 NUMBER OF SEQ ID NOS: 72 SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/10/473,126
CURRENT FILING DATE: 2003-09-26
NUMBER OF SEQ ID NOS: 1258
              TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                     APPLICANT: LEVINE, Zurit
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc_feature LOCATION: (1)..(2307) OTHER INFORMATION: any n=a,c,g,t any unknown or other
                                                ENGTH: 4533
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ENGTH: 2932
                                                                                                                                                                                                                                                                                                                                                                                                           Local
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Pred. No. 0.16;
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; LOCATION: (1)..(4533)
; OTHER INFORMATION: any n = a,c,g,t any unknown or
US-09-805-020-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-10-097-340-1/c
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US-10-072-621-2/c
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Best Local S
Matches 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 2
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CURRENT APPLICATION NUMBER: US/10/097,340 CURRENT FILING DATE: 2002-03-14 PRIOR APPLICATION NUMBER: 60/276,025 PRIOR FILING DATE: 2001-03-14 PRIOR APPLICATION NUMBER: 60/325,149
                                                                                                                                                                          TITLE OF INVENTION: Nucleic Acid Molecules and Proteins For The Identification, TITLE OF INVENTION: Assessment, Prevention, and Therapy of Ovarian Cancer FILE REFERENCE: MRI-030
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Reiner, Peter B.
APPLICANT: Connop, Bruce P.
APPLICANT: Connop, Bruce P.
APPLICANT: Pollard, Michelle
TITLE OF INVENTION: REGULATION OF AMYLOID PRECURSOR PROTEIN EXPRESSION OR J.
FILE OF INVENTION: BY MODIFICATION OF ABC TRANSPORTER EXPRESSION OR J.
FILE REFERENCE: 100103.402
CURRENT APPLICATION NUMBER: US/10/072,621
CURRENT FILING DATE: 2002-02-08
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                        APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 4643
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25; Conserv
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                                                                                                                                                                                                                                                                                Xumei ZHAO
Karen GLATT
                                                                                                                                                                                                                                                                                                                                                                               Gordon B. MILLS
Robert C. BAST, Jr.
Karen LU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sebastian HOERSCH
Shubhangi KAMATKAR
                                                                                                                                                                                                                                                                                                                                                     Rosemarie SCHMANDT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Peter VEIBY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Manjula GANNAVARAPU
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Pred. No.
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Pred. No. 0.14;
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; SEQ ID NO 1
LENGTH: 4643
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-097-340-1
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                                                                                                       RESULT 8
US-09-968-007A-459/c
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PRIOR APPLICATION NUMBER: 60/276,026
PRIOR APPLICATION NUMBER: 60/324,967
PRIOR FILING DATE: 2001-03-14
PRIOR APPLICATION NUMBER: 60/324,967
PRIOR APPLICATION NUMBER: 60/311,732
PRIOR FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: 60/325,102
PRIOR APPLICATION NUMBER: 60/325,102
PRIOR FILING DATE: 2001-09-19
PRIOR FILING DATE: 2001-09-19
PRIOR FILING DATE: 2001-09-19
                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: atp-binding cassette, sub-family b; OTHER INFORMATION: (mdr/tap), member 1 (ABCB1) gene. US-10-007-926A-258
                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 258
LENGTH: 4643
TYPE: DNA
ORGANISM: Homo sapiens
                                      Sequence 459, Application US/09968007A Publication No. US20040115625A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: BERTUCCI, FRANCOIS
APPLICANT: HOULGATTE, REMI
APPLICANT: HIRNBAUM, DANIEL
APPLICANT: NGUYEN, CATHERINE
APPLICANT: VIENS, PATRICE
APPLICANT: FERT, VINCENT
TITLE OF INVENTION: GENE EXPRESSION PROFILING OF PRIMARY BREAST CARCINOMAS
TITLE OF INVENTION: USING ARRAYS OF CANDIDATE GENES
FILE REFERENCE: 1546-R-00
CURRENT APPLICATION NUMBER: US/10/007,926A
CCURRENT FILING DATE: 2001-12-07
CCURRENT FILING DATE: 2001-12-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/254,090 PRIOR FILING DATE: 2000-12-08 NUMBER OF SEQ ID NOS: 468
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Ver. 2.1
TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Usining Signa
                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                        Local
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                                                                                                                                                                                                                                                                                                        Similarity
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                                                                                                                                                                                                GTGCTCAGCCCACGCCCCGGCGCTG 227
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                                                                                                                                                                                                                                                                                 100.0%; Score 25; DB 15; Length 4643; ilarity 100.0%; Pred. No. 0.14; Conservative 0; Mismatches 0; Indels 0
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                        Reinhard
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100.0%; Pred. No. 0.14;
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-968-007A-459
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CURRENT FILING DATE: 2001-10-02
PRIOR APPLICATION NUMBER: US/60/237,172
PRIOR FILING DATE: 2000-10-02
PRIOR PELIVATION NUMBER: US/60/237,173
PRIOR PILING DATE: 2000-10-02
PRIOR PILING DATE: 2000-10-02
PRIOR PILING DATE: 2000-10-02
PRIOR FILING DATE: 2000-10-02
PRIOR PILING DATE: 2000-10-02
PRIOR APPLICATION NUMBER: US/60/237,295
PRIOR PILING DATE: 2000-10-02
PRIOR PILING DATE: 2000-10-02
PRIOR PILING DATE: 2000-10-02
PRIOR PILING DATE: 2000-10-02
PRIOR FILING DATE: 2000-10-02
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                                                                                                                                                                                 ; TYPE: DNA; ORGANISM: Homo sapiens US-09-968-007A-747
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                                                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 2000-10-02
PRIOR APPLICATION NUMBER: US/60/237,173
PRIOR PRILING DATE: 2000-10-02
PRIOR PELLING DATE: 2000-10-02
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SEQ ID NO 459
LENGTH: 4646
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                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn version 3.0 SEQ ID NO 747
                                                                                             Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/968,007A
CURRENT FILING DATE: 2001-10-02
PRIOR APPLICATION NUMBER: US/60/237,172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Ebner Reinhard
TITLE OF INVENTION: Cancer Gene Determination
TITLE OF INVENTION: Gene Sets
FILE REFERENCE: 689290-71
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  251 GTGCTCAGCCCACGCCCCGGCGCTG 227
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                                                GTGCTCAGCCCACGCCCCGGCGCTG 25
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o. US20040115625A1
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                                                                                        100.0%;
ilarity 100.0%;
Conservative 0
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                                                                                                                Score 25; DB 1
Pred. No. 0.14;
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US-10-343-657-1/c
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                                                                                                                                                                                                                        Sequence 1, Application US/10343657 Publication No. US20040086882A1 GENERAL INFORMATION:
           APPLICANT: Roninson, Igor B.
APPLICANT: Ruth, Adam
TITLE OF INVENTION: Mutations of the MDR1 P-glycoprotein that Improve Its
TITLE OF INVENTION: Ability to Confer Resistance to Chemotherapeutic Drugs
FILE REFERENCE: 00,616-A
CURRENT APPLICATION NUMBER: US/10/343,657
CURRENT FILING DATE: 2003-10-17
PRIOR APPLICATION NUMBER: 60/222,313
PRIOR FILING DATE: 2000-08-01
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NUMBER OF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
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NAME: Zeller, Karen J.
REGISTRATION NUMBER: 37,071
REFERENCE/DOCKET NUMBER: PA-0001 US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/641,643
FILING DATE: 14-Aug-2003
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION UMBER: <Unknown>
FILING DATE: <Unknown>
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS 6.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Susan G. Stuart
Jeffrey J. Seilhamer
TITLE OF INVENTION: COMPOSITION FOR THE DETECTION OF BLOOD CELL
GENE EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: INCYTE PHARMACEUTICALS, INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Cocks, Benjamin G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 1508
                                                                                                                                                                                                                                                                                                                                                                                  251 GTGCTCAGCCCACGCCCGGCGCTG 227
SEQ ID NOS: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                           GTGCTCAGCCCACGCCCGGCGCTG 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: PALO ALTO
STATE: CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLONE: g187468
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LIBRARY: GENBANK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (650) 8.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 4646 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 25; Dilarity 100.0%; Pred. No. 0.: Conservative 0; Mismatches
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US-10-775-169-198/c
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 198
LENGTH: 4646
TYPE: DNA
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 198, Application US/10775169
Publication No. US20040175743A1
GENERAL INFORMATION:
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Best Local Simi
Matches 25;
                                                                                                                                                                                                                                                                                                                                              Sequence 1, Application US/10680516
Publication No. US20040166110A1
GENERAL INFORMATION:
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Best Local Similarity 100.0%; Pred. No. 0.14;
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SEQ ID NO 1
LENGTH: 4646
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Trepicchio, William
TITLE OF INVENTION: Method for Monitoring Drug Activities In
FILE REFERENCE: AMIO1080 (031896-013000)
CURRENT APPLICATION NUMBER: US/10/775,169
CURRENT FILING DATE: 2004-02-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Burczynski, Michael
APPLICANT: Twine, Natalie
APPLICANT: Dorner, Andrew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Wyeth
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LOCATION: (425)..(4264)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
ZIP: 60606

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff Ltd
STREET: 300 South Wacker Drive, Seventh Floor
                                                                                                                                                                                                                                                                   Roninson, Igor B
TITLE OF INVENTION: Methods and Reagents for Preparing
Using Immunoligeal Agents Specific
                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                            APPLICANT: Mechetner, Eugene
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                                                                                                                                      CITY: Chicago
STATE: Illinois
COUNTRY: USA
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100.0%; Pred. No. 0.14;
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/10/680,516 FILING DATE: 07-Oct-2003

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PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR FILING DATE: 2000-06-14
PRIOR APPLICATION NUMBER: US 60/237,054
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
SOFTWARE: PALENTIN Ver. 2.1
SEQ ID NO 3828
LENGTH: 2692
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                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Horne, Darci T.
APPLICANT: Vockley, Joseph G.
APPLICANT: Scherf, Uwe
APPLICANT: Gene Logic, Inc.
TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer
FILE REFERENCE: 44921-5928-WO
                                      Matches
                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT FILING DATE: 2001-06-14
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INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                              TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
OTHER INFORMATION: Genbank Accession No. US20020142981A1 X94563
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REGISTRATION NUMBER: 35,303
REFERENCE/DOCKET NUMBER: 95,1121
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-913-0001
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LOCATION: 4265..4669
SEQUENCE DESCRIPTION: SEQ ID NO:
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STRANDEDNESS: single
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425..4264
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87.5%; Pred. No. 42;
ative 0; Mismatches
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96.0%; Pred. No. 0.65;
tive 0; Mismatches 1;
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CURRENT APPLICATION NUMBER: US/09/968,007A
CURRENT FILING DATE: 2001-10-02
FRIOR APPLICATION NUMBER: US/60/237,172
FRIOR APPLICATION NUMBER: US/60/237,172
PRIOR PILING DATE: 2000-10-02
PRIOR APPLICATION NUMBER: US/60/237,294
PRIOR PILING DATE: 2000-10-02
PRIOR APPLICATION NUMBER: US/60/237,295
PRIOR PILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 1001
SOPTWARE: PALENTIN VERSION 3.0
SEQ ID NO 755
LENGTH: 2692
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                                                                                       Query Match
Best Local Similarity
Matches 21; Conserv
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1083 GTGCCCAGCCCCGCCGCGGCGCT 1060
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Pred. No. 42;
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Search completed: February 9, 2005, 22:26:36 Job time : 267.667 secs

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Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1: geneseqn1980s:*
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Listing first 45 summaries
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	ALIGNMENTS	ADB20870	ACF62751	AAD41243	ABS98184	ACN43504	AAQ52726	AAN70752	ADR52847	ADI31841	ADK60994	AAD38994	ABL68880	ABL68592	AAZ94738	AAQ72872	ADP54881	ADO19748	ADP18689	ABX77217	ABV74349	ABV94267	ABS76368	ABS65229	ADS88985	ADE84005	
		Adb20870 MRP1 base				•	Aaq52726 Sequence		-	Adi31841 Human CDN		Aad38994 Human mar					$\mathbf{\mu}$		Human	Abx77217 cDNA enco	Abv74349 Human ABC				U		!

Human; haematologic malignancy; multidrug resistance; MDR; SUMO-1; hypoxia inducible factor-1; small ubiquitin-like-modifier; HIF-1; lymphobid disorder; chronic lymphoproliferative disorder; lymphoma; myeloid disorder; lymphocytic leukaemia; thrombocythaemia; myeloma; angiogenic myeloid metaplasia; myeloid leukaemia; gene therapy; polycythaemia vera; hypoxia responsive element; HRE; ds. Human mdr1 gene HIF-1 binding site DNA #2. 23-SEP-2002 (first entry) AAD39009 standard; DNA; misc_binding Homo sapiens. AAD39009; 25

02-MAY-2002. 25-OCT-2001; WO200234291-A2 2001WO-US049856. Location/Qualifiers
11. 15
/*tag= a /bound_moiety= "HIF-1"

Colgan SP;

26-OCT-2000; 2000US-0243542P.

(BGHM) BRIGHAM & WOMENS HOSPITAL INC.

WPI; 2002-471427/50.

Treating a subject (at risk of) having a hematologic malignancy or multidrug resistance, e.g. lymphoma or myeloma, by administering hypoxia inducible factor 1 binding molecules or small ubiquitin-like-modifier-1 binding molecules.

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Treating a subject (at risk of) having a hematologic malignancy or multidrug resistance, e.g. lymphoma or myeloma, by administering hypoxia inducible factor 1 binding molecules or small ubiquitin-like-modifier-1
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                                                                                                                                     WPI; 2002-471427/50
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phosphorothioate backbone; ss.
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Pred. No. 4.3;
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Best Local S
Matches 25
                                                                                                                                                                                                                                                          New promoter chimera useful for in vivo or in vitro expression comprises viral promoter, intron or other transcription regulatory elements and disease-specific promoter, intron or other transcription regulatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                 King
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                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-103274/09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-MAR-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    of developing a haematologic malignancy or multidrug resistance (MD) The method involves administering hypoxia inducible factor-1 (HIF-1) binding molecules or small ubiquitin-like-modifier (SUMO)-1 binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (KING/) KING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 14; Page 43; 92pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             chimera; gene expression; viral; genotoxic; IE; hMDR1; early gene; gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; ilarity 100.0%; Conservative (
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Pred. No.
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The invention relates to a new promoter chimera for in vivo or in vitro expression that comprises: (a) a viral promoter, intron or other unit transcription regulatory sequence of any length; and (b) a unit diseasespecific promoter, intron or other unit transcription regulatory sequence of any length sensitive to genotoxic stress. The promoter chimera, and

Example 1; Page 20;

48pp; English.

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ABV75270/c
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                                                             expression that comprises: (a) a viral promoter, intron or other unit transcription regulatory sequence of any length; and (b) a unit disease-specific promoter, intron or other unit transcription regulatory sequence of any length sensitive to genotoxic stress. The promoter chimera, and vector comprising the chimera are useful for in vivo or in vitro expression. The method is useful for the generation of improved expression constructs for in vitro and in vivo expression of the present invention are more potent than either promoter chimeras of the present invention are more potent than either promoter standing alone. Sequences ABV75270-283 represent oligonucleotides used in the construction of promoter chimeras having hMDR1 disease specific DNA element and deletion of HCMV IE sequences not contributing to the activity of promoter chimera
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New promoter chimera useful for in vivo or in vitro expression comprises viral promoter, intron or other transcription regulatory elements and disease-specific promoter, intron or other transcription regulatory
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     disease-specific promoter, intron or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a new promoter chimera for in vivo or in vitro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Page 20;
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Pred. No.
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Sequence 52

BP; 9 A; 11 C; 25 G;

7 7

0 U; 0 Other;

RESULT 6

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ARESULT 5
ABV75282/c
ID ABV752
XX ABV752
XX O7-MAR
XX Promot
KW Promot
KW immedil
XX Synthe
EN WO2002
XX O7-MAR
XX O
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. Best Local S
Matches 25
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                                                                                                                                                                                                                                                                      The invention relates to a new promoter chimera for in vivo or in vitro cexpression that comprises: (a) a viral promoter, intron or other unit certanscription regulatory sequence of any length; and (b) a unit disease-ce specific promoter, intron or other unit transcription regulatory sequence of any length; sensitive to genotoxic stress. The promoter chimera, and ce vector comprising the chimera are useful for in vivo or in vitro cexpression. The method is useful for the generation of improved cexpression constructs for in vitro and in vivo expression of constructs for in vitro and in vivo expression of cexpression constructs for in vitro and in vivo expression constructs for in vitro and in vivo expression constructs for in vitro and in vivo expression of the present invention are more potent than either promoter ce thanding alone. Sequences ABV75270-283 represent oligenucleotides used in the construction of promoter chimeras having hWDR1 disease specific DNA celement and deletion of HCMV IE sequences not contributing to the activity of promoter chimera
                                                                                                               Best Loc
Matches
                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New promoter chimera useful for in vivo or in vitro expression comprises viral promoter, intron or other transcription regulatory elements and disease-specific promoter, intron or other transcription regulatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-MAR-2001; 2001US-0280211P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-MAR-2002; 2002WO-US009882
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C3 dimer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABV75282
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABV75282 standard; DNA; 56 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-103274/09
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 21; 48pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        King AC
                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (KING/) KING A C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              50 GIGCICAGCCCACGCCCCGGCGCTG
  51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 GTGCTCAGCCCACGCCCCGGCGCTG 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25;
                                                       1 GTGCTCAGCCCACGCCCCGGCGCTG 25
                                                                                                                                             Similarity
                                                                                                                                                                                                                               56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       chimera; gene expression; viral; genotoxic; IE; hMDR1; early gene; gene therapy; ss.
     GTGCTCAGCCCACGCCCCCGGCGCTG
                                                                                                                                                                                                                               B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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                                                                                                                                                                                                                               10
                                                                                                                                                                                                                               A; 11 C; 27 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  constructs creating sense oligonucleotide.
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                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                               <u>.</u>
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Pred. No.
                                                                                                                                             Score 25;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                     Mismatches
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     27
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                                                                                                                                                                       DB 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10;
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                                                                                                                                                                       Length 56;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                     Indels
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                                                                                                                     Gaps
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ABV75276/c
ID ABV75276 standard; DNA; 56 BF
XX
AC ABV75276;
XX
DT 07-MAR-2003 (first entry)
XX
C3mut3 construct creating sen
XX
Promoter; chimera; gene expre
XW
Promoter; chimera; gene; gene th
XX
C3 Synthetic.
XX
Synthetic.
XX
AC Synthetic.
XX
Synthetic.
XX
AC Synthetic.
XX
AC
RESULT 7
ABV75272/c
ID ABV752
XX
AC ABV752
XC
DT 07-MAR
XX
DE C3mut1
XX
KW Promot
KW immedi
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Best Local S
Matches 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a new promoter chimera for in vivo or in vitro expression that comprises: (a) a viral promoter, intron or other unit transcription regulatory sequence of any length; and (b) a unit disease-specific promoter, intron or other unit transcription regulatory sequence of any length sensitive to genotoxic stress. The promoter chimera, and vector comprising the chimera are useful for in vivo or in vitro expression. The method is useful for the generation of improved expression constructs for in vitro and in vivo expression of the present invention are more potent than either promoter chimeras of the present invention are more potent than either promoter standing alone. Sequences ABV75270-283 represent oligonucleotides used in the construction of promoter chimeras having hMDR1 disease specific DNA element and delection of HCMV IE sequences not contributing to the activity of promoter chimera
             Promoter; immediate
                                                                                                   C3mut1 construct creating sense oligonucleotide.
                                                                                                                                                                       07-MAR-2003
                                                                                                                                                                                                                                           ABV75272;
                                                                                                                                                                                                                                                                                                         ABV75272 standard; DNA; 56 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New promoter chimera useful for in vivo or in vitro expression comprises viral promoter, intron or other transcription regulatory elements and disease-specific promoter, intron or other transcription regulatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-MAR-2001; 2001US-0280211P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C3mut3 construct creating sense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 GTGCTCAGCCCACGCCCCGGCGCTG 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
25; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GIGCICAGCCCACGCCCGGCGCTG 27
chimera; gene expression; viral; genotoxic; IE; hMDR1;
early gene; gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 10 A; 11 C; 27 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 25;
100.0%; Pred. No.
tive 0; Mismatch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      쁌
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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RESULT 8
ABV75283
ID ABV7
XX ABV7
XX ABV7
XX Prom
XX Prom
KW Prom
KW imme
XX Synt
XX 30-N
XX 30-N
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                                                                                                                                                                                                                                                                                                                                                                                                              á
                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a new promoter chimera for in vivo or in vitro expression that comprises: (a) a vital promoter, intron or other unit transcription regulatory sequence of any length; and (b) a unit disease-specific promoter, intron or other unit transcription regulatory sequence of any length sensitive to genotoxic stress. The promoter chimera, and vector comprising the chimera are useful for in vivo or in vitro expression. The method is useful for the generation of improved expression constructs for in vitro and in vivo expression of the present invention are more potent than either promoter chimeras of the present invention are more potent than either promoter standing alone. Sequences ABV75270-283 represent oligonucleotides used in the construction of promoter chimeras having hMDR1 disease specific DNA element and deletion of HCMV IE sequences not contributing to the
                                                                                                                                      Synthetic
                                                                                                                                                                                   Promoter;
                                                                                                                                                                                                              C3 dimer and
                                                                                                                                                                                                                                             07-MAR-2003
                                                                                                                                                                                                                                                                           ABV75283;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            King
                                                                         10-OCT-2002
                                                                                                      WO200279470-A1
                                                                                                                                                                Promoter; chimera; gene expression; viral; genotoxic; IE; hMDR1; immediate early gene; gene therapy; ss.
                                                                                                                                                                                                                                                                                                        ABV75283 standard; DNA; 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New promoter chimera useful for in vivo or in vitro expression comprises viral promoter, intron or other transcription regulatory elements and disease-specific promoter, intron or other transcription regulatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 20; 48pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-103274/09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-MAR-2001; 2001US-0280211P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (KING/) KING A
                                                                                                                                                                                                                                                                                                                                                                                     27
                                                                                                                                                                                                                                                                                                                                                                                                                                                25;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                     GTGCTCAGCCCACGCCCCGGCGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                GTGCTCAGCCCACGCCCCGGCGCTG 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        56 BP; 10 A; 11 C; 27 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                              trimer constructs creating antisense oligonucleotide.
                                                                                                                                                                                                                                             (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                         ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 25; DB
Pred. No. 4.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 56;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                             0
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0

30-MAR-2001; 2001US-0280211P 29-MAR-2002; 2002WO-US009882.

(KING/) KING A C.

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RESULT 9
REVITE 9
REVITE 9
REVITE 10
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-103274/09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  King AC
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                                               New promoter chimera useful for in vivo or in vitro expression comprises viral promoter, intron or other transcription regulatory elements and disease-specific promoter, intron or other transcription regulatory
                                                                                                                                                                                                                                                                                                                                                                                                 29-MAR-2002; 2002WO-US009882.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200279470-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C3mut3 construct creating antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABV75277;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABV75277 standard; DNA; 60 BP
                                                                                                                                                                                                                                                                                                                                         30-MAR-2001; 2001US-0280211P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               immediate
                                                                                                                                                                        WPI; 2003-103274/09
                                                                                                                                                                                                                                                                                   (KING/) KING A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         chimera; gene expression; early gene; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28 C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 25; DB : Pred. No. 4.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  viral;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 10; Length 60;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          genotoxic; IE; hMDR1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0,
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The invention relates to a new promoter chimera for in vivo or in vitro cexpression that comprises: (a) a viral promoter, intron or other unit comparises: (a) a viral promoter, intron or other unit comparises: (b) a unit disease-cexpectific promoter, intron or other unit transcription regulatory sequence of any length sensitive to genotoxic stress. The promoter chimera, and cexpression. The method is useful for in vivo or in vitro cexpression. The method is useful for the generation of improved cexpression constructs for in vitro and in vivo expression of the present jene products. The HCMV IE-hMDR1 promoter chimeras of the present invention are more potent than either promoter ce standing alone. Sequences ABV75270-283 represent oligonucleotides used in the construction of promoter chimeras having hMDR1 disease specific DNA celement and deletion of HCMV IE sequences not contributing to the contivity of promoter chimera
Sequence 60 BP; 9 A; 28 C; 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 21; 48pp; English.
     G; 11 T; 0
     U; 0 Other;
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                GTGCTCAGCCCACGCCCCGGCGCTG 25
                             Conservative
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XX ABV75
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XW Immed
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                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                       C3mut1 construct creating antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                          ABV75273 standard; DNA; 60
                                                                                                                                                                                                                                              WO200279470-A1
                                                                                                                                                                                                                                                                              Promoter; chimera; gene expression; immediate early gene; gene therapy;
                                                                                                                                                                                                                             10-OCT-2002
                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                 88
                                                                                                                                                                                                                                                                                        viral; genotoxic;
                                                                                                                                                                                                                                                                                         IE; hMDR1;
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30-MAR-2001; 2001US-0280211P 29-MAR-2002; 2002WO-US009882 (KING/) KING

New promoter chimera useful for in vivo or in vitro expression comprises viral promoter, intron or other transcription regulatory elements and disease-specific promoter, intron or other transcription regulatory

WPI; 2003-103274/09.

21; 48pp; English

The invention relates to a new promoter chimera for in vivo or in vitro expression that comprises: (a) a viral promoter, intron or other unit transcription regulatory sequence of any length; and (b) a unit disease-specific promoter, intron or other unit transcription regulatory sequence of any length sensitive to genotoxic stress. The promoter chimera, and vector comprising the chimera are useful for in vivo or in vitro expression. The method is useful for the generation of improved expression constructs for in vitro and in vivo expression of the chimera perioducts. The HCMV IE-hMDR1 promoter

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RESULT 11
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                                                                                                                                                                                                                                                                                   A method of identifying (MI) a compound that modulates untranslated region-dependent expression of a vascular endothelial growth factor (VEGF) gene comprises contacting a member of a library of compounds with a cell or cell-free translation mixture containing a reporter gene operably linked to an untranslated region (UTR) of the VEGF gene, and detecting expression of the reporter gene. A compound is identified as modulator if the level of expression of the reporter gene in the presence of the compound is altered as compared to that in the absence of the compound or in the presence of a control. Compounds identified by MI are useful for treating, preventing or ameliorating cancer or its symptoms, and/or for inhibiting angiogenesis. This sequence corresponds to a
                                                                                                                                                                                      Sequence 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  chimeras of the partial standing alone. Standing alone. Standing element and delement and deleme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Identifying modulators of untranslated region-dependent expression of a VEGF gene, useful for treating cancer, comprises contacting a compound with a cell or translation mixture containing a reporter gene linked to
                                                                                                                                                                                                                                                                 therapeutic untranslated region used in the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example; SEQ ID NO 51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    VEGF gene UTR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI;
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ADR12342 standard; DNA; 424
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                                                                                                                                                                                      BP; 99
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                                                                                                                                                                                 A; 108 C; 111 G; 106 T;
        100.0%; Score 25; DB:
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cive 0; Mismatches
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Gaps
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RESULT 13
ABX94595/c
ID ABX945
XX
AC ABX945
XX
AC ABX945
XX
T 17-JUN
XX
DE Human

ABX94595

ABX94595 standard;

DNA; 633

ВP

Human MdrI 17-JUN-2003

DNA SEQ ID 3.

(first entry)

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RESULT 12
AAQ62615;

AAQ62615;

XX
AAQ62615;

XX
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AAQ62615;

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25-MAR-2003 (revised for the selection of cytostatic agents successed of the selection of cytostatic control control cytostatic control cytostatic control cytostatic control cytostatic cyt
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                                                                                                                         Matches
                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                This is a preferred fragment of the mdr-1 gene promoter, isolated from human DNA. The mdr-1 promoter and enhancer elements are inducible by cytostatic agents such as vincristine and adriamycin which are used in cancer therapy. Vectors comprising the promoter and enhancer sequences operably linked to heterologous genes coding for therapeutic agents are claimed. Using the vectors, the therapeutic agents (e.g. anti-oncogenes, interferon, IL-2, IL-6, etc.) can be expressed in cancer cells under the control of cytostatic agents. (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New mammalian expression vector useful for gene therapy, - comprising the promoter and/or the enhancer of the mdr-1 gene which is susceptible to cytostatic agents.
                                                                                                                                                                                                                                     Sequence 568 BP; 142 A; 127 C; 143 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 inducible promoter; cytostatic agent; adriamycin; vincristine;
multiple drug resistance; mdr-1; mammalian expression vector;
                                                                                                                                                  Local
      414
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                                                           -
                                                                                                                         25;
                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DELBRUECK CENT
                                                           GTGCTCAGCCCACGCCCGGCGCTG 25
GTGCTCAGCCCACGCCCCGGCGCTG 390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 6; 7pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GTGCTCAGCCCACGCCCCGGCGCTG
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                                                                                                                      Conservative
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(first entry)
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                                                                                                                                           100.0%;
                                                                                                                                                                       100.0%;
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                                                                                                                                              Score 25;
                                                                                                                      Mismatches
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                                                                                                                                                                       Length 568
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WANTE CONTROL OF THE 
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ABZ83718/c
ID ABZ83718
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention describes a novel method of determining cytosine CC methylation in a sample of genomic DNA which comprises treating the cample with hydrogen sulphite so that the cytosine is converted to uracil CC whilst 5-methylcytosine remains unchanged, amplifying sections of the DNA CC using at least 2 PCR primers and studying the base composition of both CC complementary amplified strands whereby methylation status is deduced CC from the difference in molecular weight of the two strands. The method is CC used to diagnose and/or prognose unwanted side effects of medication, CC cancer, central nervous system disorders, aggression symptoms or CC behavioural disorders, clinical, psychological and social consequence of CC brain damage, psychotic and personality disorders dementia and CC associated disorders, cardiovascular disease, malfunction, damage or CC disease of the gastrointestinal tract, breathing system, bone muscle, CC endocrine or metabolic system, injury, infection, abnormal development or sexual malfunction. This sequence represents the human MdrI DNA fragment CC amplified by the PCR primers represented in ABX94593 & ABX94594 and is CC used to illustrate the method of the invention
BXSXEXEXEXEX BXSX8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MdrI; cytosine methylation; hydrogen sulphite; diagnose; dementia; 5-methylcycosine, amplification; prognose; side effect; medication; bone; cancer; central nervous system disorder; aggression; muscle; endocrine; abnormal development; personality disorder; behavioural disorder; injury; brain damage; psychotic disorder; cardiovascular disease; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Determining cytosine methylation in a genomic DNA sample by treating hydrogen sulfite and analyzing the result, to diagnose associated conditions including cancer and brain disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2003002760-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gastrointestinal tract; sexual malfunction; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 17; 38pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-JUN-2001; 2001DE-01032212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-JUN-2002; 2002WO-DE002433.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 633
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (EPIG-) EPIGENOMICS
WO2003016500-A2
                                                                                                                                                                                            Toxicologically relevant human nucleotide sequence #877
                                                                                                                                                                                                                                                                  14-MAY-2003
                                                                                                                             Toxicologically
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     422
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GTGCTCAGCCCACGCCCCGGCGCTG 25
                                                                                                                                                                                                                                                                                                                                                                                                    standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GIGCICAGCCCACGCCCCGGCGCTG 398
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 161 A; 143 C; 165 G; 164 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                             relevant gene; toxicological response; gene;
                                                                                                                                                                                                                                                                                                                                                                                                CDNA; 1024 BP
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Pred. No. 3.6;
0; Mismatches
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CC toxicological response to an agent, which comprises comparing the cexpression profile of one or more human toxic response genes to a CC reference gene expression profile indicative of toxicity, and so CC determining the presence of a toxic response to the agent. Also CC described: (1) an array comprising one or more polynucleotides selected CC from the genes corresponding to the partial sequences given in ABZ82842 CC from the genes corresponding to the partial sequences given in ABZ82842 CC in ABZ84764, or their fragments of at least 20 nucleotides, or homologues (2) determining if a gene putatively identified to be a toxic CC response gene plays a role on toxic response pathways by determining the CC expression profile of the gene after exposure of cells or a human subject to a known toxic pharmaceutical or industrial gent, comprising: (a) CC exposing cells to an agent or isolating the test gene expression profile (comparing the test profile to the expression profile of a gene after exposure to a known toxic pharmaceutical or industrial agent; and (c) comparing the test profile to the expression profile of a gene with a similar function or comparing the test profile to the expression profile to the expression profile to the expression profile of a gene with a similar function or exposure to other known toxic compounds. The methods are useful for comparing the test profile to the agent for a cellular, organ or system level. The arrays comprising the human genes are useful for toxicological screening of drugs, pharmaceutical compounds and chemicals
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FFXGXXXXXXXX
                                                                                                                                                                                                                                                                                  AAQ62614/c
ID AAQ62
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   drugs, comprises comparing the same toxic response genes to a reference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Determining a toxicological response to an agent, useful for screening drugs, comprises comparing the expression profile of one or more human toxic response genes to a reference gene expression profile indicative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-AUG-2001; 2001US-0313080P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-AUG-2002; 2002WO-US026514
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                                                                      multiple drug resistance; mdr-1; mammalian cancer therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1024 BP; 287 A; 197 C; 260 G; 280 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (PHAS-) PHASE-1 MOLECULAR TOXICOLOGY INC.
                                                                                                                                                                                       25-MAR-2003
17-JAN-1995
                                                                                                                                                                                                                                              AAQ62614;
                                   Homo sapiens.
                                                                                                          inducible promoter; cytostatic agent; adriamycin; vincristine;
                                                                                                                                                   Human mdr-1 promoter fragment.
                                                                                                                                                                                                                                                                                  AAQ62614 standard; DNA; 1318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                                                                                251
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                       (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
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Pred. No. 3.5
0; Mismatches
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DE4238778-A1

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Search completed: February 9, 2005, 16:05:54 Job time : 247.69 secs
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                                                                                                                                                                                                          Query Match 100.0%; Score 25; DB 2; Best Local Similarity 100.0%; Pred. No. 3.4; Matches 25; Conservative 0; Mismatches 0
                                                                                                                                                                                                                                       This is a preferred fragment of the mdr-1 gene promoter, isolated from human DNA. The mdr-1 promoter and enhancer elements are inducible by cytostatic agents such as vincristine and adriamycin which are used in cancer therapy. Vectors comprising the promoter and enhancer sequences operably linked to heterologous genes coding for therapeutic agents are claimed. Using the vectors, the therapeutic agents (e.g. anti-oncogenes, interferon, II-2, II-6, etc.) can be expressed in cancer cells under the control of cytostatic agents. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                     Sequence 1318 BP; 430 A; 237 C; 277 G; 374 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                New mammalian expression vector useful for gene therapy, - comprising the promoter and/or the enhancer of the mdr-1 gene which is susceptible to cytostatic agents.
                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 2; Page 5; 7pp; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Stein U,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12-NOV-1992;
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                                                              1164 GTGCTCAGCCCACGCCCCGGCGCTG 1140
                                                                                               1 GTGCTCAGCCCACGCCCCGGCGCTG 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Walther W;
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Scoring table:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score
nucleic search, using sw
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EST: 4
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GenCore version (c) 1993 - 2005
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AW354719
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BM431190 1Duc11A08
CB431391 Duc11A08
CB423537 596986 MA
BE212025 894027B01
BF240010 BP250003B
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B0819612 1030079B0
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CD737438 4025255 1
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BU287542 603608037
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ALIGNMENTS

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JOURNAL COMMENT
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BP218711/c
                                  RESULT 2
BP348942/c
LOCUS
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                                                                                                                         472
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Euteryota; Metazoa; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 582)

Suzuki, Y., Yamashita, R., Shirota, M., Sakakibara, Y., Chiba, J., Mizushima-Sugano, J., Nakai, K. and Sugano, S. Sequence comparison of human and mouse genes reveals a homologous block structure in the promoter regions Genome Res. 14 (9), 1711-1718 (2004)

Contact: Yutaka Suzuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP218711 Sugano cDNA library, caudate nucleus Homo sapiens cDNA clone CNR05381, mRNA sequence.
BP348942 582 bp m
BP348942 Sugano cDNA library, brain
SZR02494, mRNA sequence.
                                                                                                                                                        1 GTGCTCAGCCCACGCCCCGGCGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                       Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP218711.1 GI:52091614
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                                                                                                                        GTGCTCAGCCCACGCCCCGGCGCTG
                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                             /organism="Homo sapiens"
/mol type="maNA"
/db_xref="taxon:9606"
/clone="CNRO5381"
/tissue_type="caudate nucleus"
/clone_lib="Sugano cDNA library,
                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                             100.0%; Score 25; DB 100.0%; Pred. No. 22;
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                                                                                                                  18 1 (bases 1 to 280)
18 Alzawa, K., Akahira, S., Akimura, T., Arai, A., Arakawa, T.,
18 Alzawa, K., Akahira, S., Akimura, T., Arai, A., Arakawa, T.,
18 Alzawa, K., Akahira, S., Akimura, T., Arai, A., Arakawa, T.,
18 Alzawa, K., Hanagaki, T., Hiraoka, T., Hirozane, T.,
19 Hodoyama, Y., Imotani, K., Ishii, Y., Itoh, M., Izawa, M., Kawai, J.,
19 Kojima, Y., Konno, H., Kusakabe, M., Matsuyama, T., Miyazaki, A.,
19 Nakamura, M., Nishi, K., Nomura, K., Namazki, R., Okazaki, Y.,
19 Okido, T., Owa, C., Sakai, K., Sasaki, D., Sato, K.,
19 Shibata, K., Shibata, Y., Shinagawa, A., Shiraki, T., Sogabe, Y.,
19 Shibata, K., Shibata, Y., Yasunishi, A., Yoshida, K., Yoshiki, A.,
19 Suzuki, H., Tagawa, A., Takahashi, F., Tanaka, T., Toya, T.,
19 Watahiki, A., Yamamura, T., Yasunishi, A., Yoshida, K., Yoshiki, A.,
19 Muramatsu, M. and Hayashizaki, Y.
10 Muramatsu, M. and Hayashizaki, Y.
11 REN Mouse ESTS (Aizawa, K. et al. 2000)
12 Contact: Yoshihide Hayashizaki
13 Laboratory for Genome Exploration Research Group, RIKEN Genomic Sciences Center (GSC), Yokohama Institute
11 Sciences Center (GSC), Yokohama Institute
11 Senati: 81-45-503-9216
12 Fax: 81-45-503-9216
13 Famail: Genome Exploration, Yokohama, Kanagawa 230-0045, Japan
14 Senati: 81-45-503-9216
15 Famail: Genome Exploration of the Managawa 230-0045, Japan
16 Famail: Genome Exploration of the Managawa 230-0045, Japan
17 Senatic Genome Exploration of the Managawa 230-0045, Japan
18 Senatic Genome Exploration of the Managawa 230-0045, Japan
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                                  Email: genome-res@gsc.riken.jp, URL:http://genome.gsc.riken.jp/Carninci,P., Nishiyama,Y., Westover,A., Itoh,M., Nagaoka,S., Sasaki,N., Okazaki,Y., Muramatsu,M. and Hayashizaki,Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BB608269 280 bp mRNA linear EST 06-
BB608269 RIKEN full-length enriched, 2 days pregnant adult
oviduct Mus musculus cDNA clone E230010P14 5', mRNA sequence
   Thermostabilization and thermoactivation of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims_u-tokyo.ac.jp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 (bases 1 to 582)
Suzuki,Y., Yamashita,R., Shirota,M., Sakakibara,Y., Chiba,J.,
Mizushima-Sugano,J., Nakai,K. and Sugano,S.
Sequence comparison of human and mouse genes reveals a homologous block structure in the promoter regions
Genome Res. 14 (9), 1711-1718 (2004)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus (house mouse)
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BP348942.1 GI:52278927
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/clone_lib="Sugano cDNA library, brain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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                                                                                                                                                                                                  Kanagawa 230-0045, Japan
   thermolabile enzymes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mRNA sequence
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Contact: Hyun S. Lillehoj
Contact: Hyun S. Lillehoj
Animal Parasite Diseases Laboratory
Animal and Natural Resources Institute, USDA
Animal and Natural Resources Institute, USDA
Bldg.1043, BARC-East, Beltsville, MD 20705, USA
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                                                                                                                                                                                                                                                                          clone 1GAL_80L16 5', mRNA sequence.
CD737438
CD737438.1 GI:32288287
                                                                                                                                                                        Archosauria; Aves; Neognathae; Phasianinae; Gallus.
                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Archosauria; Aves; Neognathae; Galliformes; Phasianidae;
                                                                                                                                                                                                                           Gallus gallus
                                                                                                                                                                                                                                           Gallus gallus (chicken)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     trehalose and its application for the synthesis of full length CDNA. Proc. Natl. Acad. Sci. U.S.A. 95 (2), 520-524 (1998)
Itoh, M., Kitsunai, T., Akiyama, Y., Shibata, K., Izawa, M., Kawai, J., Tomaru, Y., Carninci, P., Shibata, Y., Ozawa, Y., Muramatsu, M., Okazaki, Y. and Hayashizaki, Y. and Hayashizaki, Y. Automated filtration-based high-throughput plasmid preparation system. Genome Res. 9 (5), 463-470 (1999)
Carninci, P. and Hayashizaki, Y.
High-efficiency full-length CDNA cloning. Methods Enzymol. 303, 19-44 (1999)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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adult female oviduct"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /tissue_type="oviduct"
/dev_stage="2 days pregnant adult"
/lab_host="DH10B"
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/db_xref="taxon:10090"
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Pred. No. 1.6e+03;
0; Mismatches 3;
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gallus cDNA
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Single pass sequencing. Bases called and trimmed with phred
0.00925 using options -trim_alt '' -trim_fasta. Vector identified
by cross match using options -minmatch 12 -minscore 18
plate: 80 row: L column: 16
Seq primer: ATTTAGGTGACACTATAG
Seq primer: ATTTAGGTGACACTATAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae; Gallus.

1 (bases 1 to 652)

2 (bases 1 to 652)

2 (bases 1 to 652)

3 (bases 1 to 652)

4 (bases 1 to 652)

5 (bases 1 to 652)

6 (bases 1 to 652)

7 (bases 1 to 652)

7 (bases 1 to 652)

8 (bases 1 to 652)

9 (bases 1 to 652)

9
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Gallus gallus
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603582849F1 CSEQCHN75 Gallus gallus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Simon Hubbard
Department of Biomolecular Sciences
University of Manchester Institute of Science and Technology
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                                                                                                                                                                                                                                                                                                                                                                   Tel: 01612008930
Fax: 01612360409
                                                                                                                                                                                                                                                                                                                                                                                                                                                 PO Box 88, Manchester, M60 1QD, UK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BU382879.1 GI:25890880
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                                                                                                                                                                                                                                                                                                                                        Simon. Hubbard@umist.ac.uk.
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/ceil type="Lymphocyte"
/ceil type="Lymphocyte"
/dev_stage="Adult"
/lab_host="EMDH108"
/clone_lib="1GAL - Chicken Intestinal Lymphocyte"
/clone_lib="1GAL - Chicken Intestinal Lymphocyte"
/note="Organ: Intestine; Vector: pCMV-SPORT6; Site_1
/note="Organ: Intestine; Vector: pCMV-SPORT6; Site_1
Sall; Site_2: NotI; Normalized library from chicken
infected wIth coccidia duodenum and middle gut."
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/sex="mixed"
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/db_xref="taxon:9031"
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/mol_type="mRNA"
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                                                              /organism="Gallus gallus"

/mol type="mRNA"

/strain="White Leghorn, Hisex"

/db xref="taxon:9031"

/clone="ChEST534a14"
'lab_host="DH10B"
                                                                                                                                                                                                                                                                                                location/Qualifiers
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Pred. No. 1.4e+03;
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s cDNA clone ChEST534a14 5', mRNA
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BU122038/c
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JOURNAL
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Best Local Similarity
Matches 22; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gallus gallus (chicken)
Gallus gallus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Archosauria; Aves; Neognathae; Galliformes; Phasianinae; Gallus.

1 (bases 1 to 738)

Boardman, P.E., Sanz-Ezquerro, J., Overton, I.M., I Fong, W.T., Tickle, C., Brown, W.R.A., Wilson, S.A. A Comprehensive Collection of Chicken cDNAs Curr. Biol. 12 (22), 1965-1969 (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eute
Archosauria; Aves; Neognathae; Galliformes; Phasianidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: Simon.Hubbard@umist.ac.uk.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PO Box 88, Manchester, Tel: 01612008930
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Department of Biomolecular Sciences University of Manchester Institute
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/note="Organ: trunks; Vector: pBluescript II KS(+);
/note="Organ: trunks; Vector: pBluescript II KS(+);
Site_1: EcoRI; Site_2: Not1; This normalized library was
constructed from 1 million independent clones. cDNA
synthesis was initiated using an oligo(dT) primer, using
methylated C in the first strand synthesis reaction.
Following this first strand reaction, double-stranded cDNA
was blunted, ligated to NotI adapters, digested with
EcoRI, size-selected, and cloned into the NotI and EcoRI
compatible sites of a custom modified MCS of the
pBluescript (KS+) vector. The library was normalized in 2
rounds using conditions adapted from Soares et al., PNAS
(1994) 91: 9228-9232 and Bonaldo et al., Genome Research 6
(1996): 791, except that a significantly longer
reannealing hybridization was used."
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                                                                                                                                                                                                                                                                                                      /organism="Gallus gallus"
/mol type="mRNA"
/strain="Compton Line 151"
/db xref="taxon:9031"
/clone="ChEST147n17"
/clone lib="CSEQCHL17"
/note="Organ: kidney + adrenal; Vector: pBluescript II
/note="Organ: kidney + adrenal; Vector: pBluescript II KS(+); Site_1: EccRI; Site_2: Not1; Modification of
KS(+); Site_1: EccRI; Site_2: Not1; Modification of
pBluescript II KS(+) (Stratagene) vector to accommodate
cDNA produced with the T-trimmed protocol (Construction of
uni-directionally cloned cDNA libraries from messenger RNA
for improved 3' end DNA sequencing by Glenn Fu, et al.
U.S. Patent # 6,387,624). Cut pBluescript II KS(+) with
NotI and EccRI. Ligate in double stranded adaptor
                                                                                                                                                                                                                                  /dev_stage="adult"
/lab_host="DH10B"
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1.4e+03;
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Simon Hubbard
Department of Biomolecular Sciences
University of Manchester Institute of Science and Technology
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Eukaryota; Metazoa, Chordata; Craniata; Vertebrata; Euteleostomi;
Archomauria: Aves; Neognathae; Galliformes; Phasianidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 GTGCTCAGCCCACGCCCCGGCGCTG 25
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          Similarity 88.
22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: Simon.Hubbard@umist.ac.uk.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae; Gallus.

1 (bases 1 to 816)

1 (bases 1 to 816)

Boardman, P.E., Sanz-Ezquerro, J., Overton, I.M., Burt, D.W., Bosch, E., Broard, W.T., Tickle, C., Brown, W.R.A., Wilson, S.A. and Hubbard, S.J. A Comprehensive Collection of Chicken cDNAs

Curr. Biol. 12 (22), 1965-1969 (2002)
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603608037F1 CSEQCHN55 Gallus
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                                                                                                                                                           /dev stage="adult"
//lab host="DH10B"
//clome lib="CSEQCHN55"
//clome lib="CSE
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[5'ggccgcgtgcagccccggatccgaaaaaaag]
[5'aattctttttttcggatccggggctgcacgc]"
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/mol_type="mRNA"
/strain="Compton Line 15I"
/db_xref="taxon:9031"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sex="Female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            clone="ChEST593i10"
                                         80.8%;
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Score 20.2; DB 5;
Pred. No. 1.4e+03;
0; Mismatches 3
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s cDNA clone ChEST593i10 5', mRNA
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EST.
Gallus (chicken)
Gallus gallus
Gallus gallus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutel
Eukaryota; Metazoa; Chordata; Galliformes; Phasianidae;
                                                                                                                                                                                         l Similarity
22; Conserv
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Fax: 01612360409
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Fong, W.T., Tickle, C., Brown, W.R.A., Wilson, S.A.
A Comprehensive Collection of Chicken CDNAs
Curr. Biol. 12 (22), 1965-1969 (2002)
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University of Manchester Institute of
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                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                       /notes—Torgan: Kidney + adrenal; Vector: pBluescript II
KS(+); Site 1: EcoRI, Site 2: Not1; This normalized
library was—constructed from 1 million independent clones.
cDNA synthesis was initiated using an oligo(dT) primer,
using methylated C in the first strand synthesis reaction.
Following this first strand reaction, double-stranded cDNA
was blunted, ligated to NotI adapters, digested with
ECORI, size-selected, and cloned into the NotI and ECORI
compatible sites of a custom modified MCS of the
pBluescript (KS+) vector. The library was normalized in 2
rounds using conditions adapted from Soares et al., PNAS
(1994) 91: 9228-9232 and Bonaldo et al., Genome Research 6
(1995): 791, except that a significantly longer
reannealing hybridization was used."
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/mol_type="mRNA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        clone="ChEST802e5"
|sex="Female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /mol_type="mRNA"
/strain="Compton Line 15I"
/db_xref="taxon:9031"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             clone_lib="CSEQCHN74"
                                                                                                                                                                                                         80.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                host="DH10B"
                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                           Score 20.2; DB 5;
Pred. No. 1.3e+03;
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                                                                                                                                                                                                                            Length 929;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Euteleostomi,
                                                                                                                                                                                       Gaps
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DEFINITION

BU845700 1000 bp mRNA linear EST AGENCOURT_10414348 NIH_MGC_109 Homo sapiens cDNA clone

EST 16-OCT-2002

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REFERENCE
AUTHORS
TITLE
JOURNAL
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ORGANISM
                                                                              COMMENT
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Best Local
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BU845700
BU845700.1 GI:24030141
EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cDNA Library Preparation: Rubin Laboratory
cDNA Library Preparation: Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.lnl.gov
plate: LLCM2782 row: j column: 17
High quality sequence start: 14
High quality sequence stop: 443.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                     1017
AGENCOURT 7594073 NIH MGC 70
5', mRNA sequence.
BQ219968
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NIH-MGC http://mgc.nci.nih.gov/
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                                                                                                                                                                      Homo sapiens (human)
Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; E
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens (human)
                                      Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                        Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
                                                                                                            NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
                                                                                                                                                                                                                                                                       BQ219968.1 GI:20401357
cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                                                    (bases 1 to 1017)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /tissue_type="teratocarcinoma, cell line"
/lab_host="DH10B (phage-resistant)"
/clone lib="WIH MCC 109"
/clone orary; Vector: pOTB7; Site_1: EcoRI; Site_2:
/note="Organ: orary; Vector: pOTB7; Site_1: EcoRI; Site_2:
XhoI; cDNA made by oligo-dT priming. Directionally cloned
into EcoRI/XhoI sites using the following 5; adaptor:
GGCACGAG(G). Library constructed by Ling Hong in the
laboratory of Gerald M. Rubin (University of California,
Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and
superscript II RT (Life Technologies). Note: this is a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:6579377"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 20.2; DB 5; Length 1000; Pred. No. 1.3e+03; 0; Mismatches 3; Indels 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                  52
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sapiens cDNA clone
                                                                                                                  Gene Collection (MGC)
                                                                                                                                                                                              Euteleostomi;
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MAGE:6021580
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SOURCE
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BC015458/c
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DNA Sequencing by: Agencourt Bioscience Corporation Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov plate: LLAM13227 row: i column: 05
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                                                                                                                                     Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Series: IRAK Plate: 4 Row: f Column: 17
This clone has the following problem: retained intron.
                                                                                                                                                                                                                                                                                                                                 cDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The In.A.G.E. Consortium (LLNL)
DNA Sequencing by: Sequencing Group at the Stanford Human Genome
Center, Stanford University School of Medicine, Stanford, CA 943
Web site: http://www-shgc.stanford.edu
Contact: (Dickson, Mark) mcd@paxil.stanford.edu
Dickson, M., Schmutz, J., Grimwood, J., Rodriquez, A., and Myers,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
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Contact: MGC help desk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (01-OCT-2001) National Institutes of Health, Mammalian Gene Collection (MGC), Cancer Genomics Office, National Cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Strausberg, R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/lab_host="PH108 (phage-resistant)"
/clone_lib="NIH_MGC_70"
/clone_lib="NIH_MGC_70"
/note="Organ: pancreas; Vector: pCMV-SPORT6; Site_1: NotI;
Site_2: Sall; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.1 kb. Library constructed by Life
Technologies."
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/db_xref="taxon:9606"
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                            /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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clone="IMAGE:3456552"
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Pred. No. 1.3e+03;
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/tissue_type="Cervix, carcinoma" /clone_lib="NIH_MGC_12"

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RESULT 13
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ACCESSION
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AW354719/c
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MEDLINE
PUBMED
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Best Local
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Sequence evaluation of four pooled-tissue normalized bovine cDNA libraries and construction of a gene index for cattle
                                                                                                                                                                                                                                                                                                                                                                                     FORWARD: AGGAAACAGCTATGACCAT
BACKWARD: GTTTTCCCAGTCACGACG
Plate: 9 row: M column: 16
Seq primer: TAATACGACTCACTATAGGG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: smith@email.marc.usda.gov
Single pass sequencing. Bases called and trimmed with phred
v0.980904.e. Vector identified by cross_match with the -minscore
and -minmatch 12 options.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Smith TPL USDA, ARS, US Meat Animal Research Center PO Box 166, Clay Center, NE 68933-0166, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tel: 402 762 4366
Fax: 402 762 4390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mammalia; Eut
Bovinae; Bos.
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36851 MARC 4BOV Bos taurus cI
AW354719
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Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bos taurus
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                                                                                                                                  Similarity
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                                                                                                                                                                                               /tissue_type="pooled"
/lab_host="DH10B"
/lab_host="DH10B"
/clone_lib="MARC 4BOV"
/note="Vector: pCMV SPORT6; Site 1: Not1; Site 2
Library made from pooled tissue from day 20 and embryos."
                                                                                                                                                                                                                                                                                                  organism="Bos taurus"
/mol_type="mRNA"
/db_xref="taxon:9913"
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/note="Vector: pC
                                                                                                                                  76.8%;
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Pred. No. 3.7e
0; Mismatches
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                                                                                                                                                DB 2;
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AV589422
3', mRNA
AV589422
            and Sugimoto, Y.

Establishment of a high throughput EST sequencing system using coolv(A) tail-removed cDNA libraries and determination of 36,000
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                                                                             Takasuga, A., Hirotsune, S., Itoh, R., Jitohzono, A., Suzuki, H., Aso, H.
                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae; Bovinae; Bos.
                                                                                                                                                              Bos taurus
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EST.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Animal Genetics Division
Shirakawa Institute of Animal Genetics
Odakura, Nishigo, Nishi-shirakawa, Fuku
Tel: 81-248-25-5641
Fax: 81-248-25-5725
Nucleic
                                                                                                                                                                              Bos taurus (cow)
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This clone was obtained from a polyA-deleted cDNA library.
Location/Qualifiers
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AV663420 Bos taurus brain
3', mRNA sequence.
AV663420 .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae;
Bovinae; Bos.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bos taurus
                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: kazusugi@cocoa.ocn.ne.jp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and Sugimoto,Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AV663420.1 GI:9922450
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Yoshikazu Sugimoto
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                                                                                             (bases 1 to 452)
                                                                                                                                                                                                                                               mRNA sequence.
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                                                                                                                                                                                                                                                             Bos taurus brain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /clone_lib="Bos taurus brain fetus"
/note="Vector: pZL1; Site_1: Sal1; Site_2: Not1; Poly A
was deleted from a Not1 sīte"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /tissue_type="brain"
/dev_stage="fetus"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Bos taurus"
/mol_type="mRNA"
/db_xref="taxon:9913"
/clone="E1BR036E05"
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CDNA clone E1BR006D03
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ne E1BR036E05
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CK846580/c
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Animal Genetics Division
Shirakawa Institute of Animal Genetics
Odakura, Nishigo, Nishi-shirakawa, Fukushima 961-8061, Japan
Tel: 81-248-25-5641
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21570554
11713328
                                                                                                                                                                                                                                                                                                                                                                                     1 (bases 1 to 485)
Smith, T.P.L., Grosse, W.M., Freking, B.A., Roberts, A.J., Stone, R.T., Casas, E., Wray, J.E., White, J., Cho, J., Fahrenkrug, S.C., Bennett, G.L., Heaton, M.P., Laegreid, W.W., Rohrer, G.A., Chitko-McKown, C.G., Pertea, G., Holt, I., Karamycheva, S., Liang, F., Quackenbush, J. and Keele, J.W.

Sequence evaluation of four pooled-tissue normalized bovine cDNA libraries and construction of a gene index for cattle Genome Res. 11 (4), 626-630 (2001)
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This clone was obtained from a polyA-deleted cDNA library.
Location/Qualifiers
                                                                                                                                                                 Contact: Smith TPL
USDA, ARS, US Meat Animal Research Center
PO Box 166, Clay Center, NE 68933-0166, USA
Tel: 402 762 4366
Fax: 402 762 4390
Email: smith@email.marc.usda.gov
Single pass sequencing. Bases called with phred v0.020425.c and strimmed with the aid of the trim_alt option. Vector identified with the side of the trim_alt option.
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Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae;
Bovinae; Bos.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CK846580 mRNA linear
969088 MARC 4BOV Bos taurus cDNA 3', mRNA sequence.
CK846580
CK846580.1 GI:45207234
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                                                                                  cross match v0.990329.
Plate: 74 row: H column: 8
Seq primer: GTAATACGACTCACTATAGGG.
Location/Qualifiers
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/mol_type="mRNA"
/db_xrefe="taxon:9913"
/clone="E1BR006D03"
/clone="E1BR006D03"
/tisque_type="brain"
/dev_stage="fetus"
/lab_host="DH10B"
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/note="Vector: pZL1; Site_1: Sal1; Site_2: Not1; Poly A
was deleted from a Not1 site"
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/organism="Bos taurus"
/mol_type="mRNA"
/db_xref="taxon:9913"
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87.5%; Pred. No. 3.5e+03;
tive 0; Mismatches 3; Indels
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/lab_host="DH10B"
/clone_lib="MARC 4BOV"
/note="Vector: pCMV SPORT6; Site_1: NotI; Site_2: SalI; ...
Library made from pooled tissue from day 20 and day 40 embryos."
                                                                                                                               76.8%; Score 19.2; DB 7;
87.5%; Pred. No. 3.5e+03;
tive 0; Mismatches 3;
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Maximum Match 100%
Listing first 45 summaries
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1: gb_ba:*
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Copyright (c) 1993 - 2005 Compugen Ltd.
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AX822109

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AX597862 Sequence
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AX707915 Sequence
AC002457 Homo sapi
AC147446 Gallus ga
AC121699 Rattus no
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RESULT 2 AX504325/c LOCUS DEFINITION ACCESSION VERSION KEYWORDS SOURCE ORGANISM	Query Mat. Best Loca Matches Qy	REFERENCE AUTHORS TITLE JOURNAL FEATURES SOURCE	RESULT 1 AX504308 LOCUS DEFINITION ACCESSION VERSION VERSION KEYWORDS SOURCE ORGANISM	20 19 C 22 19 C 22 19 C 22 19 C 28 19 C 28 19 C 30 19 C 31 19 C 31 19 C 31 19 C 31 19 C 34 19 C 41 19 C 42 19 C 42 19 C 43 19 C 44 19 C 45 19
AX504325 25 bp DNA Sequence 28 from Patent WO0234291. AX504325.1 GI:23386137 Homo sapiens (human)	ch 100.0%; Score 25; DB 6; L 1 Similarity 100.0%; Pred. No. 0.67; 25; Conservative 0; Mismatches 0; 1 CCAGCATCTCCACGAAGGCAGAGTT 25	P. ions and methods for treating hemalizing resistance QTUG TESTATE HOSPITAL, INC. (US Location/Qualifiers 125 /organism="Homo sapiens"	AX504308 Sequence 11 from Patent WO0234291. AX504308 AX504308.1 GI:23386126 Homo sapiens (human) Homo sapiens (human) Eukaryota; Metazoa; Chordata; Craniata; Ve. Mammalia; Eutheria; Primates; Catarrhini; Ve.	8 79.2 272301 2 AC133673 8 79.2 349926 1 BX571660 2 76.8 59032 10 BX005046 2 76.8 6769 9 AE016815 8 6769 9 AE016815 2 76.8 121813 2 AC147587 2 76.8 129814 9 AC11664 2 76.8 130460 5 AC098806 2 76.8 148619 2 AC151162 2 76.8 148619 2 AC151162 2 76.8 168194 2 AC151462 2 76.8 181004 9 CNS01DTY 2 76.8 189228 2 AC15116 2 76.8 194313 5 BX323807 2 76.8 194313 5 BX323807 2 76.8 203839 2 AC0151268 2 76.8 203839 2 AC117692 2 76.8 203839 2 AC103892 2 76.8 221430 2 AC098629 2 76.8 230661 2 AC132770 2 76.8 230661 2 AC132770 2 76.8 230661 2 AC132072 2 76.8 346542 2 AC115307 2 76.8 346542 2 AC120727 3 76.8 346542 2 AC120727 3 76.8 346542 2 AC120727
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Isolation and sequence of the promoter region
multidrug-resistance (P-glycoprotein) gene
J. Biol. Chem. 262 (36), 17432-17436 (1987)
                                                                                                                                                                                                                                                                           Draft entry and computer-readable by I.B.Roninson, 27-OCT-1989.
                                                                                                                                                                                                                                                                                                                                        Original source text: Human multidrug
                                                                                                                                                                                                                                                                                                                                                                                        Genomic organization of the human multidrug resistance (MDR1) gene and origin of P-glycoproteins
J. Biol. Chem. 265 (1), 506-514 (1990)
                                                                                                                                                                                                                                                                                                                                                                                                                                        Chen, C.J., Clark, D., Ueda, K., Pastan, I., Gottesman, M.M. Roninson, I.B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human P-glycoprotein
M29423 J05168 M18754
M29423.1 GI:187472
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Patent: WO 0234291-A 28 02-MAY-2002;
THE BRIGHAM AND WOMEN'S HOSPITAL, INC.
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Mammalia; Eutheria;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Homo sapiens
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/gene="PGY1"
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                                                                                                                        /gene="PGY1"
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
/map="7q21"
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PUBMED REFERENCE

AUTHORS

JOURNAL MEDLINE TITLE

FEATURES

This STS was developed from sequence determined by anolinvestigator. See GenBank record: M29423 For additional information about the NHGRI chromosome 7 mapping project, http://www.nhgri.nih.gov/DIR/GTB/CHR7. Also see Genomics

mapping project, see

another

COMMENT

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                                               Buffer:
                                                                                                                           Protocol:
                                                                                                                                                                                                                                                                                                    GDB_DSEG: PGY1
Contact: Eric D. Green
Genome Technology Branch
                                                                                                                                                                                                                                                                                                                                               Human chromosome 7
Unpublished (1997)
Synonyms: PGY1
                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 976)

Bouffard,G.G., Iyer,L.M., Idol,J.R., Braden,V.V., Cunningham,A.F., Weintraub,L.A., Mohr-Tidwell,R.M., Peluso,D.C., Fulton,R.S., Leckie,M.P. and Green,E.D.
A collection of 1814 human chromosome 7-specific STSs
                                                                                                                                                                                                                         Email: egreen@nhgri.nih.gov
Primer A: AACGGAAGCCAGAACATTCC
Primer B: AGGCTTCCTGTGGCAAAGAG
                                                                                                                                                                                                                                                         National Human Génome Research
49 Convent Dr., MSC4431, Bldg.
Tel: 3014020201
Fax: 3014024735
                                                                                                                                                                                                                                                                                                                                                                                  Green, E.D.
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sWSS10 Eric
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 KCl:
Tris-HCl:
pH:
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Total Vol:
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PCR Cycles:
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Location/Qualifiers
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Sequence 136 from Patent
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
            Homo sapiens (human)
Homo sapiens
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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Sequence 1 from Patent 1
CQ806551
CQ806551.1 GI:47111933
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complement (413.
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253. .272
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253. .432
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1. .976
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Catarrhini; Hominidae;
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Berlin, K., Braun, A., Distler, J., Guetig, D., Howe, A., Mueller, J., Olek, A., Piepenbrock, C., Adorjan, P., Grabs, G., Lesche, R., Leu, E., Lewin, A., Lipscher, E., Maier, S., Model, F., Mueller, V., Otto, T., Pelet, C. and Ziebarth, H.

Methods and nucleic acids for the analysis of hematopoietic cell
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25; Conserv
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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AX457064
AX457064.1 GI:21715846
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Patent: WO 2004035803-A 1 29-APR-2004;
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AX598697
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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ilarity 100.0%;
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNI
/db_xref="taxon:9606"
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                                     Method and nucleic acids for the analysis proliferative disorder Patent: WO 03052135-A 1 26-JUN-2003; Epigenomics AG (DE)
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Sequence 1 from Patent WO03052135
AX795658
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Sequence 1 from Patent WO03044226.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                          and Nimmrich,
                                                                                                          Burger, M., Field, J.K., Genc, B., Liloglou, T., Lipscher, E.,
                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                               Homo sapiens (human)
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                                                                                                                                                                                                            AX795658.1
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            Location/Qualifiers
/organism="Homo sapiens"
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/db_xref="taxon:9606"
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/mol_type="unassigned DNJ
/db_xref="taxon:9606"
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tive 0;
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Sequence 1 from Patent WO03072821.
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Sequence 1 from Patent
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                                                                                                                       Method and nucleic acids for the analysis of a colon cell proliferactive disorder Patent: WO 03072821-A 1 04-SEP-2003;
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                                                                                                                                                             Adorjan, P., Burger, M., Rujan, T. and Schmitt, A.
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Method and nucleic acids for the analysis
prollferative disorder
Patent: EP 1340818-A 1 03-SEP-2003;
                                                                                                          Epigenomics AG (DE)
                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/db_xref="taxon:9606"
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Submitted (04-APR-1991) N. Kioka, Laboratory of Biochemistry, of Agricultural Chemistry, Kyoto University, Kitashirakawa Oiwake-cho, Sakyo-ku Kyoto 606, JAPAN See Jpn. J. Cancer Res. 80:1127-1132(1989) for overlapping
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 (bases 1 to 2932)
Kioka,N., Yamano,Y., Komano,T. and Ueda,K.
Kioka,N., Yamano,Y., Komano,T. and Ueda,K.
Heat-shock responsive elements in the induction of the multidrug
resistance gene (MDR1)
FEBS_Lett. 301 (1), 37-40 (1992)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     P-glycoprotein gene (MDR1) cDNA from human adrenal: normal P-glycoprotein carries Gly185 with an altered pattern of multidrug
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Kioka,N., Tsubota,J., Kakehi,Y., Komano,T., Gottesman,M.M.,
Pastan,I. and Ueda,K.
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X58723 X59732
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                                                                                                                                                  /gene="MDR1"
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/note="heat shock responsive element"
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KEYWORDS
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Location/Qualifiers
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 683 from Patent W003013534. AX706985
                                            Heinrich,G. and Kerb,R. Methods for treatment of cancer using Patent: WO 03013536-A 683 20-FEB-2003; Epidauros Biotechnologie AG (DE) Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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and is derived by analysis of the total score distribution.
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3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*

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5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*

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Copyright (c) 1993 - 2005 Compugen
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US-09-252-991A-13251
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US-08-615-170-20
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US-08-949-016-15083
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US-09-949-016-211849
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RESULT US-09-0 ; Seque ; Pate ; Pate ; GENE ; APP ; TIT		C 45		43						c 37					c 32		c 30		28
RESULT 1 US-09-949-016-189127 ¡ Sequence 189127, Applicat ¡ Patent No. 6812339 ¡ GENERAL INFORMATION: ¡ APPLICANT: VENTER, J. Cr ¡ TITLE OF INVENTION: POLY ¡ TITLE OF INVENTION: WIT		17								17									
189127 127, App 812339 RMATION: VENTER, VENTION:		68.0	68.0	68.0		•	•	68.0	68.0	68.0	68.0	68.0	68.0	68.8	68.8	69.6	69.6		71.2
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SULT 1 -09-949-016-189127 Sequence 189127, Application US/09949016 Sequence 18912339 GENERAL INFORMATION: APPLICANT: VENTER, J. Craig et al. APPLICANT: VENTER, J. Craig et al. TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS (ALIGNMENTS	US-09-123-708-3	US-08-365-486A-20	US-09-949-016-3696	US-09-949-016-1116	US-08-880-342-18	US-08-365-486A-18	US-09-016-434-1477	US-09-489-039A-290	US-09-902-540-3354	US-09-489-039A-419	US-09-621-976-2241	US-09-489-039A-295	US-08-984-709A-49	US-09-489-039A-294	US-09-902-540-1273	US-09-902-540-5803	US-09-949-016-17362	US-09-949-016-17434
ASSOCIATED OF DETECTION AND		Sequence :		Sequence		Sequence		Sequence				Sequence				Sequence			
USES THEREOF		3, Appli	•	3696, Ap	1116, Ap	18, App1	18, Appi	1477, Ap	290, App	3354, Ap	419, App	2241, Ap	295, App	Tddw '65	294, App	12/3, AP	5803, AD	1/364, A	17434, A

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Sequence 189128, Application US/09949016

Patent No. 6812339

GENERAL INFORMATION:

APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION A FILE REFERENCE: CL001307

CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR FILING DATE: 2000-10-20

PRIOR FILING DATE: 2000-10-3

PRIOR APPLICATION NUMBER: 60/231,768

PRIOR APPLICATION NUMBER: 60/231,768

PRIOR APPLICATION NUMBER: 60/231,768
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR PILING DATE: 2000-10-20
PRIOR PPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
VUMBER OF SED ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.
SEQ ID NO 189127
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; ORGANISM: Human
US-09-949-016-189127
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, ORGANISM: Pseudomonas aeruginosa
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                                   SEQ ID NO 13079
LENGTH: 2028
TYPE: DNA
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GENERAL INFORMATION:
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SEQ ID NO 17153
LENCTH: 76401
TYPE: DNA
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NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 189128
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Best Local Similarity
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES
FILE REFERENCE: CLO01307
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 00/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR PRILING DATE: 2000-10-03
PRIOR PILING DATE: 2000-10-03
PRIOR PILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
                                                                                                          PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
                                                                                                                                                                 TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
                                                                                                                                                                                                                                        APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                            NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 601
TYPE: DNA
ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                                                                                                       28802
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Pred. No. 1.3e+02;
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Pred. No. 6
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US-09-252-991A-12708
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                                                                                                                                                           SEQ ID NO 13351
LENGTH: 3906
TYPE: DNA
ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 13251, Application US/09252991A Patent No. 6551795
GENERAL INFORMATION:
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SEQ ID NO 12708
LENGTH: 3525
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Patent No. 6551795
GENERAL INFORMATION:
APPLICANT: MARC J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
                                                                      Matches
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Best Local
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Best Local Similarity
Matches 21; Conserv
                                                                                        Query Match
Best Local (
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CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/252,991A CURRENT FILING DATE: 1999-02-18 PRIOR APPLICATION NUMBER: US 60/074,788 PRIOR FILING DATE: 1998-02-18
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PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
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                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
 2218
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                                                                                        Similarity
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Similarity 84.0%;
                                  CCAGCATCTCCACGAAGGCAGAGTT 25
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                                                                      Conservative
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                                                                                        74.4%;
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84.0%;
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; Pred. No. 95;
0; Mismatches
                                                               Score 18.6; DB 4;
Pred. No. 1e+02;
0; Mismatches 4;
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Pred. No. 1e
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2194
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                                                                      Gaps
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APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHIANS IN KNOWN GENES ASSOCIATED

FILE REFERENCE: CL001307

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 05/241,755

PRIOR FILING DATE: 2000-10-20

PRIOR FILING DATE: 2000-10-20

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR FILING DATE: 2000-10-03

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR FILING DATE: 2000-09-08

NUMBER OF SEQ ID NOS: 207012

SOFTWARE: F886SEQ for Windows Version 4.0

1.ENOTH: 601
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; ORGANISM: Human
US-09-949-016-164913
                  APPLICATION: 435
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/01526
PILING DATE: 06-FEB-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/191,493
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Best Local Similarity 8...
20; Conservative
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Patent No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: STEWART, Alexand:
APPLICANT: LARKIN, Sarah B.
TITLE OF INVENTION: DTEF-1
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PALENLIN Rel-Base #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                      ZIP: 94105-1493
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Townsend and Townsend Khourie and Crew STREET: Steuart Street Tower, One Market Plaza CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20, Application US/08615170
>: 5776776
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AGENT INFORMATION: Heslin, James M.
                                                                                                                                                                                                                                                                                                                                                                                                                        California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FARRANCE, Iain K.G. HALL, Deborah E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MAR, Janet H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORDAHL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AZAKIE, Anthony
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Application US/09949016
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DTEF-1 ISOFORMS AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Alexandre F.R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 18.2; DB 4
Pred. No. 1.2e+02
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 4; Length 601;
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; MOLECULE TYPE:
US-08-615-170-20
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Best Local Similarity
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                          CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/01526
FILING DATE: 06-FEB-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/191,493
FILING DATE: 04-FEB-1994
CLASSIFICATION: 435
CLASSIFICATION: 435
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                              TELEFAX: (415) 326-242
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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TELEPHONE: (415) 326-2400
TELEFAX: (415) 326-2422
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/615,170
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 94105-1493
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: DT NUMBER OF SEQUENCES: 3 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Heslin, James M.
REGISTRATION NUMBER: 29,541
REFERENCE/DOCKET NUMBER: 23
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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                                                            ILECOMPUTE: (415)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 1894 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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                  ENGTH:
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: 1897 base pairs nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LARKIN, Sarah B.
/ENTION: DTEF-1 ISOFORMS AND USES THEREOF
EQUENCES: 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MAR, Janet
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORDAHL, Charles
AZAKIE, Anthony
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FARRANCE, Iain K.G.
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                                                                                        (415) 326-2400
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                                                                       326-2422
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87.0%;
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Pred. No. 1.4e+02;
                                                                                                                              23070-053120
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RESULT 11
US-08-387-942C-1
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PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4640
                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 1, Application US/08387942C Patent No. 5939289
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 72.8%;
Best Local Similarity 87.0%;
Matches 20; Conservative
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
               ZIP: 22042
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: ERTESVAG, HEI
APPLICANT: VALLA, SVEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 2087
TYPE: DNA
ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   atent No.
                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                     STREET: P.U.LU.CH
                                                                                                                                                                                                                                                                                   ITLE OF INVENTION: DNA COMPOUNDS COMPRISING SEQUENCES ITLE OF INVENTION: ENCODING MANNURONAN C-5-EPIMERASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Match 72.8%;
Local Similarity 87.0%;
APPLICATION NUMBER:
                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                        ADDRESSEE: BIRCH, STREET: P.O.BOX 747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1057 CGGCACCTCCATGAAGGCAGAGT 1035
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20; Conservative
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                                                                                                                                                   USA
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                                                                                                                                                                                                                                                                                                                  LARSEN, BJORN
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US/08/387,942C
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Pred. No. 1.4e+02;
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Pred. No. 1.4e+02;
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                                    Query Match
Best Local Similarity 87.0
                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CLOO1307
CURRENT APPLICATION UNMERER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
                                                                                                                                                              SEQ ID NO 16382
LENGTH: 36103
                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 16382, Ap
Patent No. 6812339
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Best Local Similarity
                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
                                                                                                                                                                                                               PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/241,755 PRIOR FILING DATE: 2000-10-20
                                                                                                                                                                                              SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 703-205-8050
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 12588 base pairs
                                                                                                                                               TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
NAME/KEY:
LOCATION:
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NAME/KEY:
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NAME/KEY:
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ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REGISTRATION NUMBER: 28,977
REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-205-8000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:

NAME: MURPHY JR, GERALD M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 12588 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
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1 CCAGCATCTCCACGAAGGCAGAG 23
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6702..9695
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290..1951
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9973..12588
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2227..6438
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                                                               72.8%;
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87.0%;
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                                                  0; Mismatches
                                                                 Score 18.2; DB 4; Pred. No. 2.1e+02;
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Pred. No. 1.8e+02;
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                                                                              Length 36103;
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                                                  Indels
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16613 CCAGCACCTCCACGAAAGCTGAG 16591

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GENERAL INFORMATION:

APPLICANT: VENTER, J. Craig et al.

APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

FILE REFERENCE: CL001307

CURRENT APPLICATION NUMBER: US/09/349,016

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-10-03

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR FILING DATE: 2000-09-08

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR FILING DATE: 2000-09-08

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR FILING DATE: 2000-109-08

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR FILING DATE: 2000-109-08

PRIOR FILING DATE: 2000-109-08

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR FILING DATE: 2000-109-08

PRIOR FILING DATE: 2000-109-08

PRIOR FILING DATE: 2000-199-08

PRIOR FILING DATE: 200
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Patent NO. 681233

GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF FILE REPERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/231,768
PRIOR APPLICATION NUMBER: 60/231,768
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 15083
LENGTH: 50850
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Best Local
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Query Match 72.8%;
Best Local Similarity 87.0%;
Matches 20; Conservative
                                                                                                                                                          FEATURE:
NAME/KEY: misc feature
LOCATION: (1).T. (50850)
OTHER INFORMATION: n = A,T,C or G
1-09-949-016-15084
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ORGANISM: Human
FEATURE:
NAME/KEY: misc feature
LOCATION: (1)...(50850)
OTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
ORGANISM: Human
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Pred. No. 2.2e+02;
0; Mismatches 3;
           0
           Score 18.2; DB 4;
Pred. No. 2.2e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 50850;
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                   Indels
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; NAME/KEY: misc_feature

; LOCATION: (1)...(50850)

; OTHER INFORWATION: n =

US-09-949-016-15085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-949-016-15085
Search completed: February 9, 2005, 17:11:16
Job time : 80.592 secs
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APPLICANT: VENTER, J. Craig et al.

APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

FILE REFERENCE: CL001307

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR FILING DATE: 2000-10-20

PRIOR FILING DATE: 2000-10-20

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR FILING DATE: 2000-10-03

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR FILING DATE: 2000-09-08

PRIOR FILING DATE: 2000-09-08

PRIOR FILING DATE: 2000-09-08
                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 15085
LENGTH: 50850
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 15085, Application US/09949016 Patent No. 6812339
                                                                                                                                                                Query Match
Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
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                                                                                        512 CAGACTCTCCACAAAGGCAGAGT 534
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                                                                                                                                                                        Conservative
                                                                                                                                                                                               72.8%;
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                                                                                                                                                                                                 Score 18.2; DB 4;
Pred. No. 2.2e+02;
                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                       DB 4;
                                                                                                                                                                                                                     Length 50850;
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Minimum DB seg length: 0
Maximum DB seg length: 200000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Title:
Perfect score:
Published Applications NA:*

1: /cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*

2: /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*

3: /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*

4: /cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*

4: /cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*

6: /cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*

7: /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*

8: /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*

10: /cgn2_6/ptodata/1/pubpna/US09E_PUBCOMB.seq:*

11: /cgn2_6/ptodata/1/pubpna/US09E_PUBCOMB.seq:*

12: /cgn2_6/ptodata/1/pubpna/US09E_PUBCOMB.seq:*

13: /cgn2_6/ptodata/1/pubpna/US10B_PUBCOMB.seq:*

14: /cgn2_6/ptodata/1/pubpna/US10B_PUBCOMB.seq:*

16: /cgn2_6/ptodata/1/pubpna/US10B_PUBCOMB.seq:*

17: /cgn2_6/ptodata/1/pubpna/US10B_PUBCOMB.seq:*

18: /cgn2_6/ptodata/1/pubpna/US10E_PUBCOMB.seq:*

19: /cgn2_6/ptodata/1/pubpna/US10E_PUBCOMB.seq:*

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19: /cgn2_6/ptodata/1/pubpna/US10E_PUBCOMB.seq:*

20: /cgn2_6/ptodata/1/pubpna/US10E_PUB.seq:*

20: /cgn2_6/ptodata/1/pubpna/US10E_PUB.seq:*

20: /cgn2_6/ptodata/1/pubpna/US10E_PUB.seq:*
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Maximum Match 100%
Listing first 45 summaries
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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110 8 7 6 5 110 110	Result No.
1866 1.6.66 1.6.66	Score
100.0 100.0 76.8 76.8 74.4 74.4 74.4 74.4	Query
1021 2932 561 619 2797 2914 2963 2963 2963	Query Match Length
17 18 18 17 17 17 17	DB
US-10-321-039-24 US-10-473-116-37 US-10-477-701-25185 US-10-425-115-59217 US-09-908-975-9939 US-10-336-603A-7 US-10-336-603A-7 US-10-444-575-5 US-10-40-464-125 US-10-336-603A-11 US-09-964-824A-234	ID
Sequence 24, Appl Sequence 31, Appl Sequence 25185, A Sequence 9397, Appl Sequence 9939, Appl Sequence 7, Appli Sequence 9, Appli Sequence 9, Appli Sequence 15, Appli Sequence 11, Appl Sequence 214, App	Description

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Gaps

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	US-10-369-493-45219	-10-027-632-	-10-027-632-	US-10-027-632-120970	-10-027-632-	-10-027-632-	US-10-027-632-120970	-09-876-143-314	6-143	2-674	-10-719	-10-719-	φ	US-10-152-319A-2092	ņ	US-10-719-993-13683	US-10-719-993-2097	US-10-719-993-2047	US-10-087-192-1798	US-09-805-020-27	US-10-101-510-504	US-10-087-192-1799	US-10-159-563-289	US-09-805-020-24	US-09-805-020-26	US-10-437-963-75819	US-09-805-020-25	US-10-027-632-30739	10-027	US-10-027-632-230104	US-10-027-632-230104	-094-71	US-10-336-603A-5	US-09-880-107-1654
	Sequence 45219, A	e 120972,	12097	ø	Sequence 120972,	Φ		e 314,	226,	773,	6787,	108,	e 109,	e 2092,	e 115	e 13683	2097, A	2047,	1798,	27, AI	ó	1799	æ		Sequence 26, Appl	e 758	-	35	30739,			٠.	5,	1654

ALIGNMENTS

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CURRENT APPLICATION NUMBER: US/10/321,039
CURRENT FILING DATE: 2002-12-17
PRIOR APPLICATION NUMBER: 09/998,157
PRIOR FILING DATE: 2001-11-30
PRIOR FILING DATE: 2001-10-19
PRIOR APPLICATION NUMBER: 60/360,489
PRIOR APPLICATION NUMBER: 60/360,489
PRIOR APPLICATION NUMBER: 60/360,489
PRIOR FILING DATE: 2001-10-19
NUMBER OF SEQ ID NOS: 759
SOFTWARE: PATENTIN PATENTIN OF SEQ ID NOS: 759
SOFTWARE: PATENTIN VETSION 3.2
SEQ ID NO 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 1
US-10-321-039-24
; Sequence 24, Application US/10321039
; Publication No. US20040014067A1
; GENERAL INFORMATION:
                                                                                                                       US-10-321-039-24
Query Match 100.0%; Score 25; DB 17; Length 1021; Best Local Similarity 100.0%; Pred. No. 0.076; Matches 25; Conservative 0; Mismatches 0; Indels 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Kurensky, David
TITLE OF INVENTION: Amplification Methods and Compositions
FILE REFERENCE: FORS-06960
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Lyamichev, Victor APPLICANT: Lukowiak, Andrew APPLICANT: Jarvis, Nancy
                                                                                                                                              NAME/KEY: misc feature
LOCATION: (561)...(561)
OTHER INFORMATION: n can be t or c.
                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                     FEATURE:
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                                                                                                                                                                                                                                                                                                                        1021
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US-10-767-701-25185

; Sequence 25.185, Application US/10767701

; Publication No. US20040172684A1

; GENERAL INFORMATION:

APPLICANT: Kovalic, David K.

; APPLICANT: Zhou, Yihua

; APPLICANT: Cao, Yongwei

; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With

; TITLE OF INVENTION: Plants and Uses Thereof For Plant Improvement

; FILE REFERENCE: 38-21(535)B

; CURRENT APPLICATION NUMBER: US/10/767,701

; CURRENT FILING DATE: 2004-01-29

; NUMBER OF SEQ ID NOS: 63128

; SEQ ID NO 25185
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US-10-473-126-37/c
US-10-473-126-37/c
; Sequence 37, Application US/10473126
; Publication No. US20040234973A1
; GENERAL INFORMATION:
; APPLICANT: Epigenomics AG
; TITLE OF INVENTION: Methods and nucleic acids for the analysis of hematopoietic cell
; TITLE OF INVENTION: proliferative disorders
                                                                                                              RESULT 4
US-10-425-115-59217
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                                                         Sequence 59217, Application US/10425115 Publication No. US20040214272A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                  Best Local
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CURRENT FILING DATE: 2003-09-26
NUMBER OF SEQ ID NOS: 1258
SEQ ID NO 37
                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                   Query Match
                     APPLICANT: La Rosa, Thomas APPLICANT: Kovalic, David
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Sorghum bicolor
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ORGANISM: Homo Sapiens
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zhou,
               La Rosa, Thomas J. Kovalic, David K.
                                                                                                                                                                                                                                                                Conservative
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                                                                                                                                                                                                                                                                              76.8%;
87.5%;
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                                                                                                                                                                                                                                                                                  Score 19.2;
Pred. No. 39;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-908-975-9939/c
                                                            Sequence 7, Application US/10336603A
Publication No. US20040072997A1
GENERAL INFORMATION:
APPLICANT: Alsobrook et al.
TITLE OF INVENTION: THERAPEUTIC POLYPEPTIDES,
FILE REFERENCE: Z1402-533A
CURRENT APPLICATION NUMBER: US/10/336,603A
CURRENT FILING DATE: 2003-01-03
CURRENT FILING DATE: 2003-01-03
CURRENT FILING DATE: 2003-01-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 60/221,607
PRIOR FILING DATE: 2000-07-28
NUMBER OF SEQ ID NOS: 32337
SOFTWARE: Patentin version 3.0
SEQ ID NO 9939
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 9939, Application US/09908975
Publication No. US20030165843A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants FILE REFERENCE: 38-21(5322)B FILE REFERENCE: 38-21(5322)B CURRENT APPLICATION NUMBER: US/10/425,115 CURRENT FILING DATE: 2003-04-28 NUMBER OF SEQ ID NOS: 369326 SEQ ID NO 59217 LENGTH: 619
                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
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Best Local Similarity
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Best Local Similarity
PRIOR APPLICATION NUMBER: 09/746,491
PRIOR FILING DATE: 2000-12-20
PRIOR APPLICATION NUMBER: 10/055,569
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: 36688-0005
CURRENT APPLICATION NUMBER: US/09/908,975
CURRENT FILING DATE: 2001-07-20
PRIOR APPLICATION NUMBER: US 60/287,724
PRIOR FILING DATE: 2001-05-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: SHOSHAN, Avi
APPLICANT: WASSERMAN, Alon
APPLICANT: MINTZ, BII
APPLICANT: MINTZ, LIST
APPLICANT: FAIGLER, Simchon
TITLE OF INVENTION: OLIGONUCLEOTIDE LIBRARY FOR DETECTING RNA TRANSCRIPTS AND SPLICE N
TITLE OF INVENTION: THAT POPULATE A TRANSCRIPTOME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Homo sapiens
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ORGANISM: Zea mays
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          561 CAGCATCTGCACGGAGGCAAAGTT 584
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2 CAGCATCTCCACGAAGGCAGAGTT 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 18.6;
Pred. No. 76;
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Pred. No. 39
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                                                                                                                                 NUCLEIC ACIDS ENCODING SAME, AND METHODS
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NUMBER OF SEQ ID NOS: 169
SOFTWARE: CURASeqList version 0.
SEQ ID NO 7
LENGTH: 2797
LENGTH: 2797
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (130)..(2658)
US-10-336-603A-7
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SOFTWARE: CuraSeqList version 0.1
SEQ ID NO 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 9, Application US/10336603A Publication No. US20040072997A1 GENERAL INFORMATION: APPLICANT: Alsobrook et al.
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                                                                                                                                                                                                                                                                                Sequence 5, Application US/10444575 Publication No. US20030232374A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
          APPLICANT: Kuchel, George A

APPLICANT: Zhu, Qing
TITLE OF INVENTION: Compositions and Methods Relating to Detrusor Estrogen-Regulated
TITLE OF INVENTION: Protein (DERP)
FILE REFERENCE: UCT-0035
CURRENT APPLICATION NUMBER: US/10/444,575
CURRENT FILING DATE: 2003-05-22
PRIOR APPLICATION NUMBER: US 60/382,830
PRIOR FILING DATE: 2002-05-23
NUMBER OF SEQ ID NOS: 47
SOFTWARE: Patentin version 3.2

SEQ ID NO 5
                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 2914
TYPE: DNA
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (130)..(2775)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: THERAPEUTIC POLYPEPTIDES, NUCLEIC ACIDS ENCODING SAME, AND METHOD FILE REFERENCE: 21402-533A
CURRENT APPLICATION NUMBER: US/10/336,603A
CURRENT FILING DATE: 2003-01-03
PRIOR APPLICATION NUMBER: 09/746,491
PRIOR APPLICATION NUMBER: 09/746,491
PRIOR PRIOR DATE: 2000-12-20
PRIOR APPLICATION NUMBER: 10/055,569
PRIOR PRIOR DATE: 2001-10-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -10-336-603A-9
                                                                                                                                                                                                                          APPLICANT: University of Connecticut Health Center APPLICANT: Kuchel, George A
LENGTH: 2963
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1 Similarity 84.0%;
21; Conservative
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Pred. No. 74;
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Pred. No. 74;
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APPLICANT: CHERRINGTON, JULIE
APPLICANT: SCHILLING, JIM
ITILE OF INVENTION: NOVEL BIOMARKERS OF TYROSINE KINASE INHIBITOR EXPOSURE
ITITLE OF INVENTION: AND ACTIVITY IN MAMMALS
FILE REFERENCE: 038602/1592
CURRENT APPLICATION NUMBER: US/10/440,464
CURRENT FILING DATE: 2003-05-19
PRIOR APPLICATION NUMBER: 60/380,872
PRIOR FILING DATE: 2002-05-17
PRIOR FILING DATE: 2003-02-24
PRIOR FILING DATE: 2003-02-24
PRIOR FILING DATE: 2003-02-24
NUMBER OF SEQ ID NOS: 185
SOPTWARE: Patentin Ver. 2.1
SEQ ID NO 125
LEEGTH: 2963
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DATABASE ENTRY DATE: 2003-04-07
RELEVANT RESIDUES: (1)..(2963)
US-10-444-575-5
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Publication No. US20040018528A1
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Best Local Similarity
                                                                         Matches
                                                                                        Query Match
Best Local Similarity
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AUTHORS: Tobe, T., Saguchi, K., Hashimoto, K., Miura, N.H., Tomita, M.,
AUTHORS: Li, F., Wang, Y., Minoshima, S., and Shimizu, N.
TITLE: Mapping of human inter-alpha-trypsin inhibitor family heavy
TITLE: chain-related protein gene (ITIHL1) to human chromosome 3p21-p14
JOURNAL: Cytogenet. Cell Genet.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: DEPRIMO, SAMUEL APPLICANT: O'FARRELL, ANN APPLICANT: MORIMOTO, ALYS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JOURNAL: Cyc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PAGES: 296-298
DATE: 1995
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                                                                                                                                                            TYPE: DNA ORGANISM: Homo sapiens
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1321 CCAGCTTCTCCAGGAAGGCATAGCT
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                       1 CCAGCATCTCCACGAAGGCAGAGTT 25
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                                                                       Conservative
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Pred. No. 74;
                                                                       Score 18.6; D
Pred. No. 74;
0; Mismatches
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       1297
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RESULT 10
US-10-336-603A-11/c
US-10-36-603A-11/c
; Sequence 11, Application US/10336603A
; Publication No. US20040072997A1
; GENERAL INFORMATION:

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US-09-880-107-1654/c
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; ORGANISM: Homo sapiens
US-09-964-824A-234
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                                                                                                             Sequence 1654, Application US/09880107 Patent No. US20020142981A1 GENERAL INFORMATION:
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SEQ ID NO 234
LENGTH: 3058
                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local (
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APPLICANT: HOrrigan, Stephen
TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signaturitle OF INVENTION: Sets
FILE REFERENCE: 689290-73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 234,
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                                                  APPLICANT: Horne, Darci T.
APPLICANT: Vockley, Josep
APPLICANT: Scherf, Uwe
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                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/964,824A CURRENT FILING DATE: 2001-09-27 PRIOR APPLICATION NUMBER: US/60/236,033 PRIOR FILING DATE: 2000-09-28 PRIOR APPLICATION NUMBER: US/60/236,032 PRIOR FILING DATE: 2000-09-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT FILING DATE: 2003-01-03
PRIOR APPLICATION NUMBER: 09/746,491
PRIOR FILING DATE: 2000-12-20
PRIOR APPLICATION NUMBER: 10/055,569
PRIOR FILING DATE: 2001-10-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US/60/236,028 PRIOR FILING DATE: 2000-09-28
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SOFTWARE: CuraSeqList version 0.1
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CURRENT APPLICATION NUMBER: US/10/336,603A
CURRENT FILING DATE: 2003-01-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 583
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: THERAPEUTIC POLYPEPTIDES, NUCLEIC ACIDS ENCODING SAME, AND METHOL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 2968
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
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LOCATION: (130)..(2829)
     REFERENCE:
                         OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                       Similarity
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Gene Logic, Inc.
VENTION: Gene Expression Profiles in Liver Cancer
NCE: 44921-5028-WO
                                                     Vockley, Joseph G
Scherf, Uwe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Application US/09964824A
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Pred. No. 74;
0; Mismatches
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Pred. No. 7
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; ORGANISM: Homo sapiens FEATURE: ; NAME/KEY: CDS ; LOCATION: (130)..(2919) US-10-336-603A-5
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; OTHER INFORMATION: Genbank Accession No. US-09-880-107-1654
                                                                                                                                                                                                US-10-367-094-71
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                                                                                                                                    ; Sequence 71, Application US/10367094 ; Publication No. US20040170982A1 ; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Ver. 2.1
SQ ID NO 1654
LENGTH: 3058
TYPE: DNA
ORGANISM: Homo mapiens
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Best Local S
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APPLICANT: David W. Morris
APPLICANT: Marc Malandro
ITITLE OF INVENTION: Novel Therapeutic Targets in Cancer
FILE REFERENCE: 529452001500
CURRENT APPLICATION NUMBER: US/10/367,094
CURRENT FILING DATE: 2003-02-14
NUMBER OF SEQ ID NOS: 203
                                                                                                                                                                                                                                                                                                                                                      Matches
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 09/746,491
PRIOR FILING DATE: 2000-12-20
PRIOR APPLICATION NUMBER: 10/055,569
PRIOR FILING DATE: 2001-10-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: 21402-533A
CURRENT APPLICATION NUMBER: US/10/336,603A
CURRENT FILING DATE: 2003-01-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT FILING DATE: 2001-06-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Alsobrook et al.
TITLE OF INVENTION: THERAPEUTIC POLYPEPTIDES, NUCLEIC ACIDS ENCODING SAME, AND METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 60/237,054
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
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84.0%;
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Pred. No. 74
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; Pred. No. 74;
0; Mismatches
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SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 71

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Sequence 230104/c

US-10-027-632-230104/c

Sequence 230104, Application US/10027632

Publication No. US20020198371A1

GENERAL INFORMATION:

APPLICANT: Wang, David G.

TITLE OF INVENTION: Identification and Mapping of Single Nucleotide

TITLE OF INVENTION: Polymorphisms in the Human Genome

FILE REFERENCE: 108827.129

CURRENT APPLICATION NUMBER: US/10/027,632

CURRENT FILING DATE: 2002-04-30

PRIOR APPLICATION NUMBER: US 60/218,006

PRIOR PILING DATE: 2000-07-12

PRIOR PILING DATE: 2000-07-12

PRIOR PILING DATE: 2000-03-29

PRIOR PILING DATE: 2000-03-29

PRIOR APPLICATION NUMBER: US 60/193,483

PRIOR APPLICATION NUMBER: US 60/195,218

PRIOR APPLICATION NUMBER: US 60/195,363

PRIOR APPLICATION NUMBER: US 60/185,218

PRIOR APPLICATION NUMBER: US 60/167,363

PRIOR APPLICATION NUMBER: US 60/167,363

PRIOR APPLICATION NUMBER: US 60/167,363

PRIOR PILING DATE: 1999-09-28

PRIOR APPLICATION NUMBER: US 60/167,363

PRIOR PILING DATE: 1999-09-28

PRIOR PILING DATE: 1999-09-28

PRIOR PILING DATE: 1999-09-09

NUMBER OF SEQ ID NOS: 325720

SOFTWARE: PASTSEQ for Windows Version 4.0

1 ENGTH- 4644
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; TYPE: DNA
; ORGANISM: Human
; 0-10-027-632-230104
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ORGANISM: Mus musculus
PEATURE:
NAME/KEY: misc_feature
LCCATION: (1)...(684187)
OTHER INFORMATION: n = A,T,C or G
US-10-367-094-71
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                                                                                                                                                                                                   Query Match 72.8%; Score 18.2; DB 13; Best Local Similarity 87.0%; Pred. No. 1.2e+02; Matches 20; Conservative 0; Mismatches 3;
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PT Treating a subject (at tiek of maring a nemacologic militaring promitted resistance, e.g. lymphoma or myeloma, by administering	WPI; 2002-471427/50.	PI Colgan SP; XX	PA (BGHM) BRIGHAM & WOMENS HOSPITAL INC.			PD 02-MAY-2002.	WO200234291-A2.	FH Key Location/Qualifiers FT modified_base 125 FT /*tag= a FT /*mod_base= OTHER FT /note= "Phosphorothioate backbone"	OS Homo sapiens. OS Synthetic. XX	KW Human; haematologic malignancy; multidrug resistance; MDR; SUMO-1 KW hypoxia inducible factor-1; small ubiquitin-1ike-modifier; HIF-1; Wallymphorid disorder; chronic lymphoproliferative disorder; lymphomak KW myeloid disorder; lymphocytic leukaemia; thrombocythaemia; myelom KW angiogenic myeloid metaplasia; myeloid leukaemia; gene therapy; KW polycythaemia vera; hypoxia responsive element; HRE; antisense; KW phosphorothioate backbone; ss.	OT 23-SEP-2002 (first entry)	AD39000;	RESULT 1 AAD39000 B AAD39000 standard; DNA; 25 BP.	ALIGNMENTS	45 18.2 72.8 444 4 AAI79965	43 18.6 74.4 5793 13 ACN42391	41 18.6 74.4 5783 13 ACN42392	40 18.6 74.4 5672 13 ACN42394	38 18.6 74.4 4042 10 ADBU7449 39 18.6 74.4 4226 12 ADQ64433	37 18.6 74.4 3906 11 ABD14647	35 18.6 74.4 3489 8 ACC59960 36 18.6 74.4 3525 11 ABD14104	33 18.6 74.4 3058 10 ADD49032 34 18.6 74.4 3144 10 ADG75752	31 18.6 74.4 3058 6 ABL41972 32 18.6 74.4 3058 6 ABN95156	29 18.6 74.4 2968 10 ADD49038 30 18.6 74.4 3058 6 ABL67200	27 18.6 74.4 2963 12 ADHI7126 28 18.6 74.4 2963 12 ADI19055	25 18.6 74.4 2960 13 ACN37942 26 18.6 74.4 2963 6 ABL41971	23 18.6 74.4 2905 12 ADJ72005 24 18.6 74.4 2914 10 ADD49036	21 18.6 74.4 2797 10 ADD749034 22 18.6 74.4 2834 12 ADJ72006	
cering hypoxia										SUMO-1; HIF-1; ymphoma; myeloma; rapy; ense;					965 Human pol	2391 Human dia inuation (3 of	2392 Human dia	2394 Human dia 2393 Human dia	4433 Novel hum	4647 Pseudomon	960 Human PMM 4104 Pseudomon	5752 Human pro	Abn95156 Gene #165	200 Thyroid c	9055 Human int	7942 IUMOUL-AB 971 DNA seque	2.7	2006 Human PMM	_

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Description

SUMMARIES

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Aad39000 Human mdr Aad39010 Human mdr Abz09897 Human 5° Adb53945 MDR1 geno Ade84005 5° regula Ads88985 Human ABC

ADB53945 ADE84005 ADS88985 ABS98184 AAD41243 ACF62751

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ADB20870 ADB87959 ADB967959 ADB9692133 ABN37191 AD044565 ADC39482 ADC39482 ADC98671 ADC98676 ADC98080 ADP28076 ADP28076 ADP28076

Abs98184 Human mul
Abs98184 Human mul
Aad41243 Human MDR
Acf62751 Cancer ba
Adb20870 MRP1 base
Adb807959 Human MDR
Adb96942 Human MDR
Adb969133 Human MDR
Abn37191 Human spl
Add44565 AI480570e
Adc39482 Novel hum
Ade09661 Novel bum
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Ade09661 Novel bum
Ade09661 Human sec
Adp28080 Human sec
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Crime method involves administering hypoxia inducible factor-1 (HIF-1) binding molecules or small ubiquitin-like-modifier (SUMO)-1 binding consecutes or HIF-1-SUMO-1 complex modulators. mdr1-hypoxia responsive celement (HRE) binding molecules or antisense nucleic acid molecules and SUMO-1 binding molecules or antisense molecules are useful for treating a celement (HRE) binding molecules or antisense molecules are useful for treating a subject having or at risk of developing haematologic malignancy or MDR (e.g. a lymphoid disorder or a myeloid disorder). The lymphoid disorders cinclude lymphocytic leukaemia or chronic lymphoproliferative disorders ce.g. lymphoma or chronic lymphoid leukaemia. The myeloid disorders include chronic or acute myeloid leukaemia, e.g. angiogenic ce.g. lymphomatolasia, essential thrombocythaemia or polycythaemia vera. The invention is used in gene therapy. The present sequence is an antisense colligo targetted to HRE in the mdr1 gene (mdr1-HRE) to inhibit its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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                    Treating a subject (at risk of) have multidrug resistance, e.g. lymphoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; haematologic malignancy; multidrug resistance; MDR; SUMO-1; hypoxia inducible factor-1; small ubiquitin-like-modifier; HIF-1; lymphoid disorder; chronic lymphoproliferative disorder; lymphoma; myeloid disorder; lymphocytic leukaemia; thrombocythaemia; myeloma; angiogenic myeloid metaplasia; myeloid leukaemia; gene therapy;
                                                                                                              WPI; 2002-471427/50
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                                                                                                                                                                    Colgan
                                                                                                                                                                                                                                                                                     26-OCT-2000;
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                                                                                                                                                                                                                             WOMENS HOSPITAL INC
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                 risk of) having a hematologic malignancy oe.g. lymphoma or myeloma, by administering
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ng hypoxia
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RESULT 3
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ID ABZ098
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DT 16-JAN
XX Human;
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Best Local S
Matches 25
                                                                            Detecting and differentiating between hematopoietic cell proliferative disorders, comprises contacting a target nucleic acid with a reagent that distinguishes between methylated and non-methylated CpG dinucleotides.
                                                                                                                                                                                       Berlin K, Braun A, Distler J,
Olek A, Piepenbrock C, Adorjan
Lewin A, Lipscher E, Maier S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  element (HRE) binding molecules or antisense nucleic acid molecules and SUMO-1 binding molecules or antisense molecules are useful for treating subject having or at risk of developing haematologic malignancy or MDR (e.g. a lymphoid disorder or a myeloid disorder). The lymphoid disorders include lymphocytic leukaemia or chronic lymphoproliferative disorders e.g. lymphoma, myeloma or chronic lymphoid leukaemia. The myeloid disorders include chronic or acute myeloid leukaemia, e.g. angiogenic myeloid metaplasia, essential thrombocythaemia or polycythaemia vera. The invention is used in gene therapy. The present sequence is human mdrl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a method of treating a subject having or at of developing a haematologic malignancy or multidrug resistance (MDR). The method involves administering hypoxia inducible factor-1 (HIF-1) binding molecules or small ubiquitin-like-modifier (SUMO)-1 binding molecules or HIF-1-SUMO-1 complex modulators mdrl-hypoxia responsive
                                                                                                                                                                         Schwope I,
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                                                                                                                                                                                                                                                                                                                 26-MAR-2002;
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ilarity 100.0%;
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1 P, Grabs
Model F,
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differentiating

invention describes a method for detecting and invention describes a method for detecting and

disorders

Claim 38;

SEQ

ID NO

37; 117pp; English.

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RESULT 4

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ID ADB539

XX ADB539

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XX O4-DEC

XX Colon

KW Cytost

XX Uniden

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        disorders allowing for improved and informed treatment of patients
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between gene or nucleic

The invention

colon cell proliferative disorders associated with at least its regulatory regions. The method comprises contacting a ta acid in a biological sample obtained from the contacting a tax acid in a biological sample obtained from the contacting at a scid in a biological sample obtained from the contacting at a scid in a biological sample obtained from the contacting at a scid in a biological sample obtained from the contacting at a scid in a biological sample obtained from the contacting and differential colors.

differentiating

The invention relates to a method of detecting and differentiating between lymphoid cell proliferative disorders associated with at least one gene and/or their regulatory regions in a subject by contacting a target nucleic acid in a biological sample obtained from the subject with at least one reagent or series of reagents that distinguish between methylated and non-methylated CpG dinucleotides within the target nucleic acid. The genes and/or their regulatory regions are preferably selected from MDR1, CSNX2B, EGR4, AR, CDK4, RB2, CDC25A, GPJL beta, MYOD1, CDH3, MYCL1, ELKI, ABL1, APC, BCL2, CDH1, CDKN1A, CDKN1B, CDKN2B, FOS, GSTP1, HIC-1, MGMT, MLH1, MOS, MYC, PTEN, RBL2, TGFBR2, TP73, CDKNIC,

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RESULT 5
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XX Ad; ly
KW diffus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           methylated CpG dinucleotide; single nucleotide polymorphism; diffuse large B-cell lymphoma; mantle cell lymphoma; chronic lymphocytic leukemia; small lymphocytic lymphoma; follicular lymphoma; diagnosis; prognosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5' regulatory region of human gene MDR1.
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                                                                                                                                                                                                                                                                                                                                                               Detecting and differentiating between lymphoid cell proliferative disorders comprises contacting a target nucleic acid with at least reagent that distinguishes between methylated and non-methylated Cr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ds; lymphoid cell proliferative disorder; methylation;
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                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 1; 448pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Burger M,
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28-DEC-2001;
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2001DE-01064501.
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Matches
The invention relates to a novel method for predicting the responsiveness of a subject with a cell proliferative disorder of the breast tissues to a therapy comprising analysing the methylation pattern of a target nucleic acid by contacting at least one of the target nucleic acids in a biological sample obtained from the subject prior to or during treatment. The method of the invention has cytostatic activity, and may have a use in gene therapy. The set of oligonucleotides comprising at least two of the oligomers are useful for detecting the cytosine methylation state and/or single nucleotide polymorphisms (SNPs) within the sequences. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GSK3beta, ESR1, APAF1, BAK1, BAX or HOXA5. Oligomers, peptide nucleic acid (PNA)-oligomers and/or isolated nucleic acids based on the sequences of the genes are useful for detecting the methylation state of all the cpG dinucleotides within one or more the sequences, or their complements, for determining the cytosine methylation state and or single nucleotide polymorphisms (SNPs), and for differentiating at least two of the medical conditions such as diffuse large B-cell lymphoma, mantle cell lymphoma, chronic lymphocytic leukemia, small lymphocytic lymphoma and follicular lymphoma. They are also useful for detecting of a predisposition to, differentiation between subclasses, diagnosis, prognosis, treating and/or menitoring of lymphoid cell proliferative disorder. This sequence menitored for and/or regulatory region from one of the above
                                                                                                                                                                                                               Predicting responsiveness of a subject with breast cell disorder, useful for treating or differentiating breast proliferative disorders comprises analyzing methylation
                                                                                                                                                                                                                                                                                                                                                                                              01-OCT-2002; 2002DE-01045779.
07-JAN-2003; 2003DE-01000096.
17-APR-2003; 2003DE-01017955.
                                                                                                                                                                   Example 2;
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                                                                                                                                                                                                                                                                                                                                   Foekens J,
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Rujan T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gene
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Schmitt
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pattern of a
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Marx A;
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RESULT 7
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KW cytochrome P450 A2; CYP4501A2; cytochrome P450 02E; CYP4502E1; LTF;
KW adrenergic receptor beta1; ADBR1; arryl hydrocarbon; AHR; MRP3; NR112;
KW aryl hydrocarbon receptor nuclear translocator; ARNT; cathepsin S; CTSS;
KW cyclooxgenase 2; COX2; diazepam binding inhibitor; DBI; haematological;
KW cyclooxgenase 2; EPHX2; 5-lipoxygenase activating protein; FLAP;
My plutathione-5-transferase 12; GST12; histamine-N-methyl transferase;
KW HNMT; kallikrein 2; KLK2; nicotinamide-N-methyl transferase; NNMT;
KW MADPH quinone oxidoreductase 2; NQO2; sulfotransferase thermolabile; STM;
KW MADPH quinone oxidoreductase 214; UDD-glucuronosyl transferase 2B7;
WP-glucuronosyl transferase; UGT2B15; urokinase receptor; uPA;
KW UDP-glucuronosyl transferase; UGT2B15; urokinase receptor;
KW multidrug resistance 1; lactotransferin; orphan nuclear receptor;
KW multidrug resistance associated protein 3; cancer; prostate;
MY acetylcholine muscarinic receptor; CHMR1; CHMR2; CHMR3; CHMR4; CHMR5;
KW altered drug metabolism; cardiovascular function; colorectal tumour;
KW altered associated protein 3; cancer; prostate;
acetylcholine muscarinic receptor; upa;
WW central nervous system; pulmonary; immunological; SNP;
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Matches 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CCAGCATCTCCACGAAGGCAGAGTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CCAGCATCTCCACGAAGGCAGAGTT 1570
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 resistance
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gene promoter B and exons 1-3 sequence.
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nucleotide

polymorphism.

WO200257410-A2

25-JUL-2002

28-NOV-2001; 2001WO-US044838.

28-NOV-2000; 2000US-00724389

(DNAS-) DNA SCI LAB

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2002-698522/75

Isolated nucleic acid molecules having polymorphisms in known human e.g. cytochrome p450 and cathepsin S useful as genetic linkage marke for locating, identifying and characterizing the genes responsible f disorder-related traits. enetic linkage markers genes responsible for

432-437; 714pp; English

This invention relates to the sequence of an isolated nucleic acid molecule comprising at least one base variation from that of a known human cytochrome P450 A1 (CYP4501A1), cytochrome P450 A2 (CYP45002E1), adrenergic receptor beta1 (ADBR1),

a

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RESULT 8
ADD4124/c
ID AADD412
XX AAD412
XX Human
XX Human
XX Human
XX Human
XX Homo 8
XX P-glyc
XX W02002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; transgenic; reporter construct; CYP3A4; cytochrome transcriptional regulatory element; xenobiotic; steroid; Mp-glycoprotein; gene; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 8573 BP; 2739 A; 1422 C; 1705 G;
   Liddle C,
                                                                                                                                 01-NOV-2000;
10-MAY-2001;
                                                                                                                                                                                                                             01-NOV-2001; 2001WO-AU001407.
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                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
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                                                                    (UNSY ) UNIV SYDNEY
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          Goodwin BJ,
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                                                                                                                                 2000AU-00001161.
2001AU-00004901.
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          Robertson
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RESULT 9
ACF62751/c
ID ACF62751
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to generation of non-human transgenic animals comprising a reporter construct for producing a detectable amount of a reporter molecule operably linked to a transcriptional regulatory nucleic acid molecule from the human CYP3A4 gene. The invention also relates to transgenic animal for analysing CYP3A4 cytochrome P450 gene regulation. The non-human mammal of the invention is useful for determining whether a compound is capable of effecting the transcription of a human CYP3A4 gene. It is also useful for determining the effect of a compound such as a xenobiotic or steroid on the regulation of expression of the CYP3A4 gene in a human. The present sequence is 5' flanking region of human (p-glycoprotein) MDR1 gene. This sequence is used as a regulatory element in the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Non-human mammal for determining the effect of a compound on regulation of CYP3A4 gene expression, comprises a reporter construct for producing reporter linked to a transcriptional regulatory nucleic acid from a human reporter linked to a transcriptional regulatory nucleic acid from a human reporter linked to a transcriptional regulatory nucleic acid from a human reporter linked to a transcriptional regulatory nucleic acid from a human reporter linked to a transcriptional regulatory nucleic acid from a human reporter linked to a transcription regulation r
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                       9817
                                                                                                                                                                                                                                                                                       1 CCAGCATCTCCACGAAGGCAGAGTT
                                                                                                                                                                                                                                                                                                                                                                                                                  25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
         standard; DNA; 177380
                                                                                                                                                                                                                                       CCAGCATCTCCACGAAGGCAGAGTT 9793
                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Score 25; 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
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Cancer; CYP3A5; irinotecan; pharmaceutical; malignant glioma; cytochrome p450; subfamily IIIA; nifedipine oxidase; polypeptide 5; cytostatic; gene; ds.
                                                                                                                                                                                                                                                                                                                                                                 Cancer based on CYP3A5 related polynucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                          08-OCT-2003
The present invention describes the use of irinotecan (I) or its derivative for the preparation of a pharmaceutical composition for
                                                          New use of irinotecan for preparation of compositions for treating cancer in subject having genome with variant allele comprising cytochrome p450, subfamily IIIA, polypeptide 5 polynucleotide, termed CYP3A5.
                                                                                                                                                                                23-JUL-2001; 2001EP-00117608.
24-MAY-2002; 2002EP-00011710.
                                                                                                           WPI; 2003-268144/26.
                                                                                                                                                                                                                    23-JUL-2002; 2002WO-EP008219
                                                                                                                                                                                                                                              20-FEB-2003
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                                                                                                                                                                                                                                                                                            Unidentified
                                                                                                                                   Heinrich G,
                                                                                                                                                          (EPID-) EPIDAUROS BIOTECHNOLOGIE
                                      SEQ
                                                                                                                                   Kerb
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                                      ID NO
                                                                                                                                   æ
                                        683;
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                                      English.
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RESULT 10
ADB20870/c
ID ADB208
XX ADB208
XX ADB208
XX ININCT
DT 20-NOV
XX irinot
KW irinot
KW Lung c
KW varian
KW Varian
KW Varian
KW Varian
XX CPID-
XX (EPID-
XX Heinri
XX YAR
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XX Use of
PT Compos
PT Varian
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a cytochrome p450, subfamily IIIA (nifedipine oxidase), polypeptide 5 (CYPAS) polynucleotide (II). (I) and (II) have cytostatic activity. The therapeutic applications of (I) is improved, since it is possible to individually treat a subject with an appropriate desirvative of (I). Therefore, undesirable, harmful or toxic effects are efficiently avoided. Unnecessary and potentially harmful treatment of those subjects who do not respond to the treatment with substances (nonresponders), as well as the development of drug resistances due to suboptimal drug dosing can be avoided. ACF62200 to ACF62751 and ABM34912 to ABM35013 represent sequences used in the exemplification of the present invention
                                                                                                                                                                                                            Use of irinotecan or its derivative for preparation of a pharmaceutical composition for treating cancer in a subject having a genome with a variant allele comprising a multidrug resistance protein 1 polynucleotide.
                                                                                                                                                                                                                                                                                                                                                                                         23-JUL-2001;
24-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  irinotecan; colorectal cancer; cervical cancer; gastric cancer;
lung cancer; ovarian cancer; pancreatic cancer; malignant glioma;
variant allele; multidrug resistance protein 1; MRP1; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                          23-JUL-2002; 2002WO-EP008200
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cancer related nucleic acid SEQ ID NO:683.
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2002EP-00011710.
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Pred. No. 0.6
); Mismatches
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The present invention describes a method for the use of irinotecan (I) of its derivative for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a multidrug resistance protein 1 (MRP1) polynucleotide (II). (I) has cytostatic accivity, (I) or its derivative can be used for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject, where the subject is a human

Disclosure; SEQ

ID NO 683; 100pp; English.

or

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CCAGCATCTCCACGAAGGCAGAGTT

25

Query Match Best Local (Matches

Similarity

100.0%;

Score 25; Pred. No.

0.61; 멂

10; <u>و</u>

Length 177380;

Conservative

<u>,</u>

Mismatches

0;

Indels

0

Gaps

0

Sequence

177380

B₽;

58338 A; 32530 C; 31389

55123 T;

0 U; 0 Other;

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Best Local S
Matches 25
                         The invention relates to the novel use of irinotecan to treat a patient suffering from cancer. This involves determining if the patient has one or more variant alleles of the UGTIA1 gene, and if the patient has one or more of such variant alleles, irinotecan is administered in an increased or decreased amount in comparison to the amount that is administered without regard to the patient's alleles in the UGTIA1 gene. The invention has cytostatic activity. A composition of the invention acts as a topoisomerase I inhibitor. The method is useful for treating a patient, an animal e.g. mouse or a human, preferably African or Asian, suffering from cancer such as colorectal, cervical, gastric cancer, lung, ovarian, pancreatic cancer or malignant glioma. The present sequence is udes in the exemplification of the invention.
                                                                                                                                                                                                                                                 Use of irinotecan to treat cancer patient by determining if patient has variant alleles of UGTIA1 gene, administering increased/decreased amounts
                                                                                                                                                                                                          Disclosure; SEQ ID NO 683; 107pp; English.
                                                                                                                                                                                                                                       of irinotecan based
                                                                                                                                                                                                                                                                                                                       Heinrich
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          irinotecan; cancer; UGT1A1; cytostatic; topoisomerase I :
colorectal cancer; cervical cancer; gastric cancer; lung
ovarian cancer; pancreatic cancer; malignant glioma;
                                                                                                                                                                                                                                                                                                                                                                           23-JUL-2001;
24-MAY-2002;
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2002EP-00011710.
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Pred. No.
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RESULT 12
ADB96942/c
ID ADB969
XX ADB969
XX ADB969
XX ADB969
XX ADB969
DT 04-DEC
XX Irinot
XX Irinot
XX Irinot
XX TOP1;
XX TOP1;
XX TOP1;
XX 20-FEE
XX 23-JUL
XX 21-FEE
XX 23-JUL
XX 21-FEE
XX 21-FE
RESULT 13
ADB92133/c
ID ADB921
XX
AC ADB921
AC ADB921
XX
DT 04-DEC
XX
DE Human 1
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Best Local S
Matches 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to the novel use of irinotecan or its derivative for the preparation of pharmaceutical compositions for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a multidrug resistance 1 (MDR1) polynucleotide. A composition of the invention has cytostatic activity. The invention is useful for the preparation of pharmaceutical compositions for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject (preferably human, more preferably African or Asian) or a mouse. The present sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        irinotecan; colorectal cancer; cervical cancer; gastric cancer;
lung cancer; ovarian cancer; pancreatic cancer; malignant glioma;
multidrug resistance 1; MDR1; cytostatic; human; Cyp3A5; MRP1; MDR1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New use of irinotecan for preparation of pharmaceutical compositions for treating cancer in subject having genome with variant allele comprising multidrug resistance 1 polynucleotide.
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24-MAY-2002; 2002EP-00011710.
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         Human MDR1
                                                               04-DEC-2003
                                                                                                                                                                               ADB92133 standard;
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                                                                                                                                                                                  DNA; 177380
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100.0%; Pred. No. 0.0
:ive 0; Mismatches
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            SEQ ID NO:683
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RESULT 14
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Matches 25
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lung cancer; ovarian cancer; pancreatic cancer; malignant glioma;
multidrug resistance 1; MDR1; cytostatic; human; UGTIA1; MRP1; TOP1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates ro a novel use of irinotecan or its derivative for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a multidrug resistance 1 (MDR1) polynucleotide. A composition of the invention has cytostatic activity. The present sequence is used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New use of irinotecan for preparation of pharmaceutical compositions for treating cancer in subject having genome with variant allele comprising multidrug resistance 1 polynucleotide.
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24-MAY-2002; 2002EP-00011710.
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                                                                                                                                                                       Human;
splice
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          exemplification of the invention.
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                                                                                                                                                                                                                                                                                    ABN37191 standard; DNA; 60 BP
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                                                                                                                                                                                                            Human
                                                                                                                                                                                                                                    15-JUL-2002
                                                                                                                                                                                                                                                              ABN37191;
                                 28-JUL-2000; 2000US-0221607P.
02-MAY-2001; 2001US-0287724P.
                                                                                                                         WO200210449-A2
                                                                       20-JUL-2001; 2001WO-IB001903
                                                                                                07-FEB-2002.
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                                                                                                                                                                                                             spliced
                                                                                                                                                                      mouse; rat; splice transcript; detection; RNA transcript; variant; transcriptome; oligonucleotide library; ss.
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25; Conserv
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ilarity 100.0%;
Conservative (
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                                                                                                                                                                                                          transcript detection oligonucleotide SEQ ID NO:9939
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(COMP-) COMPUGEN INC

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RESULT 15
AD044565
ID AD044
XX AD044
XX AD044
XX AD1480
XX CRH;
KW GRH;
KW GRH
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                                                                                                                                                                                                                    23-OCT-2003; 2003WO-EP011792
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CRH; corticotropin-releasing hormone; antidepressant; tranquillizer; gene therapy; CNS; expressed sequence tag; EST; ds.
   Fierens
                                Peeters
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 60 BP; 16 A; 15 C; 16 G; 13 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription unit of
                                                                                               (JANC ) JANSSEN PHARM NV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AI480570expressed sequence tag (EST)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes oligonucleotide libraries for detecting messenger RNAs that populate a (sub-)transcriptome, where the (sub-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shoshan A,
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   FLP;
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                                Goehlmann HWH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                Kass
                                su,
                                Steckler THW
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Diagnosing a corticotropin-releasing hormone (CRH)-induced gene expression profile in an individual comprises determining polypeptides and polynucleotides that mediate the endocrine response to CRH in the
                                                                                                                                            Claim 1; SEQ ID NO 31; 84pp; English.
                                                                                                                                                              sample
                                                                                                                                                                                                        WPI; 2004-376160/35.
GENBANK; AI848545.
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The invention relates to diagnosing a corticotropin-releasing hormone (CRH)-induced gene expression profile in an individual. The method involves determining the level of gene transcription of a gene comprising a nucleic acid sequence selected from 28 nucleotide sequences fully defined in the specification or determining the amount of at least one protein that modulates CRH signaling in the sample, where the protein comprises 13 amino acid sequences given in the specification. The methods and composition are useful for diagnosing, treating and/or preventing CRH metabolism-related disorders, such as CRH-induced depression or stress. The polymucleotide is used as a marker of CRH signaling in a cell. These composition and methods may also be used in identifying compounds that modulate CRH-induced depression and stress. The present sequence represents a specific example of a gene fragment that is an important mediator of CRH-induced changes in the CNS.

Sequence 354 BP; 107 A; 71 C; 68 G; 108 T; 0 U; 0 Other;

밁 Ś Query Match Best Local S Matches 21 1 CCAGCATCTCCACGAAGGCAGAGTT 25 21; Similarity Conservative 74.48; 0; Score 18.6; DB: Pred. No. 2e+02; 0; Mismatches 12; 4 Length Indels 0, Gaps

0

rch completed: February time : 247.69 secs 262 CCAGCAGCTCCACGAAGTCAGTGAT 286 9, 2005, 16:05:56

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Title:
Perfect score:
Sequence:
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No.
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Run on:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Total number of hits satisfying chosen parameters:
                                                                                                                                                                                              0000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score
     EST: *
2: 90,0
3: 90,0
4: 90,0
5: 0
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Match
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25
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Gapop 10.0 , Gapext 1.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             34239544 seqs, 19032134700 residues
     ccagcatctccacgaaggcagagtt
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gb_est2:*
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9b_est4:*
9b_est5:*
9b_est6:*
9b_gs81:**
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     DB
  AV631826
AV626483
BZ536972
CG206154
CG206156
BZ785280
CG436517
BM257699
AL726135
BX307891
AL726201
AL726201
AL726201
AL726201
AC29563
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AV631826 AV631826
AV626483 AV626483
BZ536972 OGAHH18TC
CG206154 PUJDB60TD
CG436506 CGVCE10TV
BM257659 521522 MA
AL726135 AL726135
BX307891 BX307891
BX1726201 AL726201
AV594929 AV594929
BX299583 BX299583
CR364140 CR364140
CR3664140 CR366111-16L1
AV59482 BX299583
BX299583 BX299583
CR364140 CR364140
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ALIGNMENTS

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| Qy
Db 4 | Query Match
Best Local :
Matches 2: | ORIGIN | | actions | FEATURES | | | COMMENT | PUBMED | MEDITINE | | 7777.5 | AUTHORS | REFERENCE | ORGANISM | SOURCE | VERSION | DEFINITION | Locus | RESULT 1
AV631826 |
| 1 CCAGCATCTCCACGAAGGCAGAGTT 25

 174 CCAGCATCTTCACAAATGCAGAGTT 498 | / Match 80.8%; Score 20.2; DB 1; Length 505;
Local Similarity 88.0%; Pred. No. 4.1e+02;
1es 22; Conservative 0; Mismatches 3; Indels 0; Gaps 0; | | /mol_type="mkNA" /strain="09" /db_xref="taxon:3055" /db_xref="taxon:3055" /clone="LC1100d12_r" /clone="lb="Chlamydomonas reinhardtii 5% to 0.04% CO2" /clone="Vector: pBluescriptII SK-; Site 1: EccRI; Site_2: /note="Vector: pBluescriptII SK-; Site 1: EccRI; Site_2: /note="Vector: pBluescriptII SK-; Site 1: EccRI; CO2" /note="Vector: pBluescriptII SK-; Site 1: EccRI; Site_2: /note="Vector: pBluescriptII SK-; Site_1: EccRI; Site_2: EccRI; EccRI; Site_2: EccRI; Site_2: EccRI; E | organism="Chlamydomonas reinhardtii" | Location/Qualifiers | Yana 1532-3, Kisarazu, Chiba 292-0812, Japan
Email: asamizu@kazusa.or.jp, URL:http://www.kazusa.or.jp/en/plant/. | The First Laboratory for Plant Gene Research
Kazusa DNA Research Institute | Contact: Brika Agamizu | 11089912 | DNA KES. / (5), 305-30/ (2000) | cells of Chlamydomonas reinhardtii | nd Tabata, S. | Asamizu, E., Miura, K., Kucho, K., Inoue, Y., Fukuzawa, H., Ohyama, K., | | Chlamydomonas reinnardtii Eukaryota; Viridiplantae; Chlorophyta; Chlorophyceae; Volvocales; | mydomonas | AV631826.1 GI:10794460
EST. | AV631826 CHIAMIYUUMUHAB IGIIMIATUCII 3° CO 0.018 CO2 CHIAMIYUCHAO 1000 CCL100d12_r 5', mRNA sequence: AV631826 CHIAMIYUUMUHAB IGIIMIATUCII 3° CO 0.018 CO2 CHIAMIYUUMUHAA 1826 CHIAMIYUUMUHAA 1826 CHIAMIYUUMUHAA 1826 CHIAMIYUUMUHAA 1826 CHIAMIYUUMUHAA 1826 CHIAMIYUUMUHAA 1826 CO2 CHIAMIYUUMUHAA 1826 CHIAMIYUUMUHAA 1826 CHIAMIYUUMUHAA 1826 CHIAMIYUUMUHAA 1826 CHIAMIYUUMUHAA 1826 CO2 CHIAMIYUUMUHAA 1826 CHIAMIYUUMUHAA 1826 CHIAMIYUUMUHAA 1826 CO2 CHIAMIYUUMUHAA 1826 CO2 CHIAMIYUUMUHAA 1826 CO2 CHIAMIYUUMUHAA 1826 CHIAMIYUUMUHAA 1826 CO2 CHIAMIYUUMUHAA 182 | | |

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AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 463
                                                            Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopaida; Poales; Poaceae; PACCAD clade; Panicoideae; Andropogoneae; Zea.

1 (bases 1 to 702)

Whitelaw,C.A., Quackenbush,J., Van Aken,S., Utterback,T., Resnick,A., Fraser,C.M., Budiman,M.A., Bedell,J.A., Rohlfing,T., Citek,R.W., Nunberg,A., Robbins,D. and Lakey,N. Consortium for Maize Genomics Unpublished (2002)
Contact: Cathy Whitelaw
                                                                                                                                                                                                                                                                                                                                        OGAHH18TC ZM2_0.7_1.5_KB genomic survey sequence. BZ536972
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Asamizu, E., Miura, K., Kucho, K., Inoue, Y., F. Makamura, Y. and Tabata, S. Generation of expressed sequence tags from adapted cells of Chlamydomonas reinhardtii DNA Res. 7 (5), 305-307 (2000)
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                                                                                                                                                                                                                                                                                                                      BZ536972.1 GI:27085170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yana 1532-3, Kisarazu, Chiba 292-0812, Japan
Email: asamizu@kazusa.or.jp, URL:http://www.kazusa.or.jp/en/plant/.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Erika Asamizu
The First Laboratory for Plant Gene Research
Kazusa DNA Research Institute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chlamydomonadaceae; Chlamydomonas.
1 (bases 1 to 514)
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                            Medical Center Drive,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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/clone="LCL009h12 r"
/clone="LCL009h12 r"
/clone=11b="Chlamydomenas reinhardtii 5% to 0.04% CO2"
/clone_11b="Chlamydomenas reinhardtii 5% to 0.04% CO2"
/note="Vector: pBluescriptII SK-; Site_1: EcoRI; Site_2:
XhOI; The cDNA library was constructed from cells cultured
in a carbon stress acclimatized condition in which carbon
dioxide concentration in the bubbling gas was changed from
5% to 0.04%"
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/mol_type="mRNA"
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Pred. No. 4.1e+02;
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Zea mays
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                                                                                                                                                                                                                                                                                                                         9712 Medical Center Drive,
Tel: 301-838-5843
Fax: 301-838-0208
                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (2003)
Other_GSSs: PUJDB60TB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fax: 301-838-0208
Email: whitelaw@tigr.org
                                                                                                                                                                                                                                                                                                                                                                                    Contact: Cathy Whitelaw
                                                                                                                                                                                                                                                                                                                                                                                                                                Maize Genomics Consortium
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                                                                      Similarity
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Bennetzen,J.
CCAGCATCTGCACGGAGGCAAAGTT
                           CCAGCATCTCCACGAAGGCAGAGTT 25
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/clone_lib="ZM_0.6_1.0_KB"
/note="Vector: pCR4-TOPO; Site_1:
CoT selected genomic DNA library"
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/clone_lib="ZM2_0.7_1.5_KB"
/note="Vector: pBCSK-; Site_1
/methylation filtered genomic
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/strain="B73"
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1 (bases 1 to 852)

1 (bases 1 to, 852)

1 (bases 1, 0, 852)

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Other_GSSs: OGVGE10TV
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1 (bases 1 to 893)

whitelaw,C.A., Quackenbush,J., Van Aken,S., Utterback,T., Resnick,A., Fraser,C.M., Yuan,Y., San Miguel,F., Ma,J. and
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                                                     Maize Genomics Consortium Unpublished (2003) Contact: Cathy Whitelaw
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9712 Medical Center Drive, Rockville, MD 20850, USA
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/mol type="genomic DNA"
/strain="B73"
/db_xref="taxon:4577"
/clone="ExMSMA0475A19"
/clone="ZM/BMA0475A19"
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/clone="ExMSMA0475A19"
/clone="ExMSMA0475A19"
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/note="Vector: pBCSK: Site 1: HincII; 0.7-1.5
methylation filtered genomic DNA library"
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CGVGE10TV ZM 0.7 1.5 KB Zea m
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Class: sheared
                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD clade; Panicoideae; Andropogoneae; Zea.

1 (bases 1 to 955)

Whitelaw, C.A., Quackenbush, J., Van Aken, S., Utterback, T., Resnick, A., Fraser, C.M., Budiman, M.A., Bedell, J.A., Rohlfing, T., Citek, R.W., Nunberg, A., Robbins, D. and Lakey, N. Consortium for Maize Genomics Unpublished (2002)
Other_GSSs: OGVEENOTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 CCAGCATCTCCACGAAGGCAGAGTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fax: 301-838-0208
Email: whitelaw@tigr.org
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tel: 301-838-5843
                           1 CCAGCATCTCCACGAAGGCAGAGTT
                                                                                                                                                                                                                                                                                                                                            9712 Medical Center Drive,
Tel: 301-838-5843
Fax: 301-838-0208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zea mays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Zea mays
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                                                                                                                                                                                                                                                                                                   Class: sheared ends.
                                                                                                                                                                                                                                                                                                              Seq primer: TF
                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Cathy Whitelaw
                                                                                                                                                                                                                                                                                                                               Email: whitelaw@tigr.org
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CCAGCATCTGCACGGAGGCAAAGTT 391
CCAGCATCTGCACGGAGGCAAAGTT
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                                                              Conservative
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/clone="myMSHB2771724"
/clone="myMSHB2771724"
/clone="myMSHB2771724"
/clone="person" 0.6 1.0 KB"
/note="Vector: pCR4-TOFO; Site_1:
/note="Vector: pCR4-TOFO; Site_1:
COT selected genomic DNA library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Zea mays"
/mol_type="genomic D
/strain="B73"
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                                                                                                                                       /clone="ZMMBMa0475A19"
/clone_lib="ZM 0.7_1.5_KB"
/note="Vector: pBCSK-; Site_1: HincII; 0.7-1.5
methylation filtered genomic_DNA_library"
                                                                                                                                                                                                       db xref="taxon:4577"
                                                                                                                                                                                                                    /mol_type="genomic DNA"
/strain="B73"
                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                  organism="Zea mays"
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Pred. No. 4.5e+02;
                                                            Score 20.2; DB 9;
Pred. No. 4.5e+02;
0; Mismatches 3;
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                                                                                                                                                                                                                                                                                                                                                                                  Rockville,
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SOURCE
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                                                                                                                                               DEFINITION
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Matches 21
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BM257699/c
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MEDLINE
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                                                                                                                                                                                                                                                                       122 CCAGCACCTCCACGAAGGCTGAG 100
  Danio rerio (zebrafish)
Danio rerio
Eukaryota; Metazoa; Cho
                                                                                                                   AL726135
AL726135 Danio rerio embryonic inner ear subtracted rerio CDNA clone BNOAA083ZHO2 5', mRNA sequence.
                                                                                AL726135
AL726135.1
                                                                                                                                                                                                                                                                                                                                                         21;
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Smith, T.P.L., Grosse, W.M., Freking, B.A., Roberts, A.J., Stone, R.T., Casas, E., Wray, J.E., White, J., Cho, J., Fahrenkrug, S.C., Bennett, G.L., Heaton, M.P., Laegreid, W.W., Rohrer, G.A., Chitko-McKown, C.G., Pertea, G., Holt, I., Karamycheva, S., Liang, F., Chitko-McKown, C.G., Pertea, G., Holt, I., Karamycheva, S., Liang, F., Chackenbush, J. and Keele, J.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BACKWARD: GTTTTCCCAGTCACGACG
Plate: 125 row: D column: 5
Seq primer: ATTTAGGTGACACTATAG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Smith TPL
USDA, ARS, US Meat Animal Research Center
PO Box 166, Clay Center, NE 68933-0166, USA
Tel: 402 762 4366
Fax: 402 762 4390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: smith@email.marc.usda.gov
Single pass sequencing. Bases called and alt_trimmed with phred
v0.980904.e. Vector identified by cross_match with the -minscore 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence evaluation of four pooled-tissue normalized bovine cDNA libraries and construction of a gene index for cattle Genome Res. 11 (4), 626-630 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BM257699 550 bp mRNA linear 521522 MARC 3BOV Bos taurus cDNA 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                       /db_xréf="taxon:9913"
/tissue_type="pooled"
/tissue_type="pooled"
/lab_host="D4108"
/clone_lib="MARC_3BOV"
/clone_lib="MARC_3BOV"
/note="Vector: pCMV_sPORT6; Site_1: NotI; Site_2: SalI_Library made from pooled tissue from marrow, alveolar macrophage, ovary, fetal semitendonosus muscle, and felongissimus muscle."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Bos taurus"
/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           location/Qualifiers
Metazoa; Chordata; Craniata; Vertebrata;
                                                                                  GI:20190739
                                                                                                                                                                                                                                                                                                                                                                         79.2%;
91.3%;
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                                                                                                                                                                                                                                                                                                                                                     Score 19.8; DB 4;
Pred. No. 6.2e+02;
0; Mismatches 2
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                                                                                                                                                                                                                                                                                                                                                                                            Length 550;
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                                                                                                                                           EST 18-APR-2002
CDNA Danio
Euteleostomi;
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BX307891/c
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AUTHORS
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Best Local Similarity
Matches 21; Conserv
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                                                                                                                                                                                                                                                                                                               INRA - SCRIBE
Campus de beaulieu, I
Tel: 02.23.48.50.09
Fax: 02.23.48.50.20
                                                                                                                                                                                                                                                  Email: Yann.Guiguen@beaulieu.rennes.inra.fr
Sequence cleaned of vector, adaptator and re
at sigenasupport@jouy.inra.fr to obtain the
                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (2003)
On Apr 7, 2003 this sequence version replaced gi:29588536.
Contact: Guiguen Y
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  w
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Govoroun, M., Guiguen, Y. and Le Gac, F.
Construction and primary characterization of normalized cDNA
libraries in rainbow trout, Oncorhynchus mykiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Actinopterygii; Neopterygii; Teleostei; Euteleostei; Protacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus. 1 (bases 1 to 585)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oncorhynchus mykiss (rainbow trout)
Oncorhynchus mykiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mRNA sequence.
BX307891
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Genoscope
Genoscope - Centre National de Sequencage
2 rue Gaston Cremieux, CP 5706 - 91057 EVRY cedex - FRANCE
Email: segref@genoscope.cns.fr, Web : www.genoscope.cns.fr.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes; Cyprinidae; Danio.

1 (bases 1 to 584)
Coimbra,R., Weil,D., Brottier,P., Blanchard,S., Levi, Hardelin,J.P., Weissenbach,J. and Petit,C.
A subtracted cDNA library from the zebrafish (Danio r
                                                                                                                                                                                                                                  sequence
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                                                                                                                                                                                                               Plate: 0016
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Unpublished (2002)
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                                                                                                                                                                                          primer: M13R.
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/organism="Oncorhynchus mykiss"
/mol_type="mRNA"
/db_xref="texon:8022"
/clone="texo0:8022"
/clone="type="adipose tissue, blood, brain,
/tissue_type="adipose tissue, blood, brain,
differentiating gonads, gills, interrenal, intestine,
kidney, liver, muscle, ovary, pituitary, testis"
                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone_lib="Danio rerio embryonic cDNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /tissue_type="inner ear"
/dev_stage="embryonic"
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/mol_type="mRNA"
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                                                                                                                                                                                                               column:
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Pred. No. 6.3e+02;
0; Mismatches 2;
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                                                                                                                                                                                                                                                  repetitions. Co
he chromatogram
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AV594929/c
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AL726201 Danio rerio embryonic
rerio cDNA clone BNOAA083ZH02:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Danio rerio (zebrafish)
Danio rerio
AV594929 Bos taurus cartilage fe ElCA016F04 5', mRNA sequence. AV594929 AV594929.1 GI:9711387
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Genoscope - Centre National de Sequencage
2 rue Gaston Cremieux, CP 5706 - 91057 EVRY cedex -
Email: segref@genoscope.cns.fr, Web : www.genoscope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 bases 1 to 587)
Colmbra,R., Weil,D., Brottler,P., Blanchard,S., Levi,M.,
Hardelin,J.P., Weissenbach,J. and Petit,C.
A subtracted cDNA library from the zebrafish (Danio rerio)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AL726201.1 GI:20190805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Genoscope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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/note="Vector: pT/T3D-pac; Rainbow trout multi-tissues ./
/note="Vector: pt/T3D-pac; PIUMI, Claration ./
/note="Vector: pt/T3D-pac; PIUMI, Claration ./
/note="Vector: pt/T3D-pac; PT/T3D-
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/lab_host="DH10B"
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/tissue_type="inner ear"
/dev_stage="embryonic"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        organism="Danio rerio"
/mol_type="mRNA"
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Pred. No. 6.3e+02;
0; Mismatches 2;
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Pred. No. 6.3e+02;
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                                                                                              fetus Bos taurus cDNA
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inner ear subtracted
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                               INRA - SCRIBE
Campus de beaulieu, l
Tel: 02.23.48.50.09
Fax: 02.23.48.50.20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Yoshikazu Sugimoto
Animal Genetics Division
Shirakawa Institute of Animal Genetics
Odakura, Nishigo, Nishi-shirakawa, Fuk
Tel: 81-248-25-5641
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Establishment of a high throughput EST poly(A) tail-removed cDNA libraries and
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                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Actinopterygii; Neopterygii; Teleostei; Euteleostei; Protacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus.
                                                                                                                                                                                                                                                                                                                     EST.
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Mammalia; Eutheria;
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                                                                                                                     Unpublished (2003)
On Apr 7, 2003 this sequence version replaced gi:29580228.
                                                                                                                                                        Govoroun,M., Guiguen,Y. and Le Gac,F.
Construction and primary characterization of normalized cDNA
libraries in rainbow trout, Oncorhynchus mykiss
                                                                                                                                                                                                                                                                                  Oncorhynchus mykiss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: kazusugi@cocoa.ocn.ne.jp
                                                                                                                                                                                                                                                                                                 Oncorhynchus mykiss (rainbow trout)
                                                                                                                                                                                                                                                                                                                                    BX299583.2
                                                                                                                                                                                                                                                                                                                                                                       mRNA sequence.
Email: Yann.Guiguen@beaulleu.rennes.inra.fr
Sequence cleaned of vector, adaptator and repetitions. Contact us
                                                                                                        Contact: Guiguen Y
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/note="Vector: pZL1; Site 1: Sal1; Site
was deleted from a Not1 sIte"
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/dev_stage="fetus"
/lab_host="DH10B"
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/clone="B1CA016F04"
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/mol_type="mRNA"
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                                                                     RENNES
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Pred. No. 6.3e+02;
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                                                                                                                                                                                                                  Seq
                                                                                                                                                                                                                                       Plate: 0038,
                                                                                                                                                                                                                                                                                                                                               Campus de beaulieu, RENNES cedex, Tel: 02.23.48.50.09
Fax: 02.23.48.50.20
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Construction and primary characterization of normalized cDNA libraries in rainbow trout, Oncorhynchus mykiss Unpublished (2003)
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1 (bases 1 to 687)

Govoroun, M., Guiguen, Y. and Le Gac, F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oncorhynchus mykiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CR364140 687 bp mRNA linear EST 21-APR-2004 CR364140 AGENAE Rainbow trout normalized testis library (tcbi) Oncorhynchus mykiss cDNA clone tcbi0038c.j.24 Sprim, mRNA sequence. CR364140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N
                                                                                                                                                                                                                                                                                at sigenasupport@jouy.inra.fr to
                                                                                                                                                                                                                                                                                                      Email: Yann.Guiguen@beaulieu.rennes.inra.fr
Sequence cleaned of vector, adaptator and re
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oncorhynchus mykiss (rainbow trout)
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                                                          /organism="Oncorhynchus m
/mol type="mRNA"
/db xref="texon:8022"
/clone="tcbi0038c.j.24"
/tissue_type="testis"
                                                                                                                                                                                                                                                                                                                                                                                                                                     Guiguen Y
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/note="Vector: pT7T3D-pac; Rainbow trout multi-tissues -
/note="Vector: pT7T3D-pac; Rainbow trout multi-tissues -
normalized + 1 subtraction (tcay); Clone distribution :
AGENAE Resource centre. Francois PIUMI,
Francois.Piumi@jouy.inra.fr, INRA, CEA Radiobiologie et
Etude du genome (LREG), Domaine de Vilvert, 78352,
Jouy-en-Josss cedex, France, +33 (0) 1.34.65.28.02, +33
(0) 1.34.65.22.73"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /tissue_type="adipose tissue, blood, brain,
differentiating gonads, gills, interrenal, intest
kidney, liver, muscle, ovary, pituitary, testis"
/dev_stage="from embryos to adults"
/lab_host="DH10B"
/Clone_lib="AGENAE Rainbow trout (tcbi)"
                                                                                                                                                                                              Location/Qualifiers
                                             lab host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
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/mol_type="mRNA"
/db_xref="taxon:8022"
/clone="tcay0004b.i.10"
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91.3%;
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0; Mismatches 2
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CN069432/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AUTHORS
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646 CCAGCACCTCCACGAATGCAGAG
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                                                                                                                                                                                                                                                                                                                              Department of Biology
University of Kentucky
TH Morgan Building, Lexington, KY 40506, USA
Tel: 859 257 988
Fax: 859 257 1717
Email: srvos@uky.edu
The EST is quality trimmed at the ends with a 20 base window and quality threshold of 15 (phred quality score).Please visit http://salamander.uky.edu For any information(trace,quality files etc) regarding this EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ambystomatid salamanders
BMC Genomics 5 (1), 54 (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Putta,S., Smith,J.J., Walker,J.A., Rondet,M., Weisrock,D., Monaghan,J., Samuels,A.K., Kump,K., King,D.C., Maness,N.J., Habermann,B., Tanaka,E., Bryant,S.V., Gardiner,D.M., Parichy,D.M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ambystoma tigrinum tigrinum
Eukaryota; Metazoa; Chordata;
Amphibia; Batrachia; Caudata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Salamander_28_M15.ab1 AG Ambystoma tigrinum tigrinum
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: SR Voss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         From biomedicine to natural history research: EST resources for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and Voss, S.R.
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                                                                                        Similarity
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                                                                       Conservative
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                                                                                                                                                            /mol type="mRNA"

(sub_species="tigrinum"

/db_sref="taxon:43116"

/tissue_type="fiver, Lung, Kicgill tissues collected from me/clone_lib="AG"
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Francois.Piumi@jouy.inra.fr, INRA, CEA Radiobiologie et
Etude du genome (IREG), Domaine de Vilvert, 78352,
Jouy-en-Josas cedex, FRANCE, +33 (0) 1.34.65.28.02, +33
(0) 1.34.65.22.73"
                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                            organism="Ambystoma"
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                                                                     Score 19.8; DB 7;
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m metamorphosing larvae"
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Search completed: February Job time: 2451.85 secs

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Listing first 45 summaries
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Maximum DB seq length: 200000000
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   ORGANISM Homo sapiens
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| 4553 | 4378 | 4264 | 4264 | 4195 | 4195 | 4192 | 4186 | 4186 | 3988 | 3988 | 3988 | 3860 | 3860 | 3852 | 3852 | 3852 | 3843 | 3840 | 966 | 807 | 807 | 473 | 473 | 473 | 463 |
| Q | σ | σ | σ | σ | σ | 9 | σ | σ | 9 | σ | σ | σ | σ | 9 | ø | φ | σ | თ | ø | σ | 9 | σ | σ | δ | δ |
| CQ716151 | E02326 | AR051650 | AR051647 | AX108656 | AR399195 | AF016535 | AX108654 | AR399194 | AX024454 | AR452556 | BD190394 | AX322789 | AX322787 | AY582534 | AF537134 | AF537133 | BD171402 | AX481416 | HUMMDR1A07 | AR363343 | I08556 | CQ332256 | 567 | CQ136561 | CQ331579 |
| CQ716151 Sequence | c | 1650 Sequence | 1647 | 656 | 5 Sequ | 6535 | 4 Sequ | 4 | | AR452556 Sequence | | 2789 | 7 | 4 | 7134 | 7133 Macaca m | 1402 | 6 Sequenc | | 4 | equence | CQ332256 Sequence | | 36561 Sequenc | w |

ALIGNMENTS

| AX504309 | Avendano 25 hn DNA linear PAT 27-SEP-2002 |
|--|---|
| DEFINITION | 12 from Patent WO0234291. |
| ACCESSION | |
| VERSION | AX504309.1 GI:23386127 |
| KEYWORDS | |
| SOURCE | Homo sapiens (human) |
| ORGANISM | |
| | etazoa; Chordata; |
| | Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. |
| REFERENCE | P |
| AUTHORS | |
| TITLE | Compositions and methods for treating hematologic malignancies and |
| | multiple drug resistance |
| JOURNAL | Patent: WO 0234291-A 12 02-MAY-2002;
THE BRIGHAM AND WOMEN'S HOSPITAL, INC. (US) |
| FEATURES | Location/Qualifiers |
| source | 125 |
| | <pre>/organism="Homo sapiens" /mol_type="unassigned DNA" /db_xref="taxon:9606"</pre> |
| 1 | |
| Query Match
Best Local :
Matches 2 | <pre>Match 100.0%; Score 25; DB 6; Length 25; Local Similarity 100.0%; Pred. No. 0.059; les 25; Conservative 0; Mismatches 0; Indels 0; Gaps 0;</pre> |
| δ | 1 AGCTTCCAACCACGTGTAAATCCTA 25 |
| DЪ | 1 AGCTTCCAACCACGTGTAAATCCTA 25 |
| RESULT 2
AX504326/c | |
| Locus | AX504326 25 bp DNA linear PAT 2/-SEF-2002 |
| DEFINITION | Sequence 29 from Patent WO0234291. |
| VERSION | AX504326.1 GI:23386138 |
| SOURCE | Homo sapiens (human) |
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RESULT 4
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Kit for assaying MDR1 gene expression level

L Patent: JP 2002153274-A 3 28-MAY-2002;

KAINOS LAB INC

OS Artificial Sequence
PN JP 2002153274-A/3

PD 28-MAY-2002

17-NOV-2002

PI 7-NOV-2000 JP 2000351224

PI TAKESHI UJIIE,TSUKASA HAYASHI,MASAKO KAWASE,FUMIAKI UEJI PC

C12N15/09,C12Q1/68,C12N15/00

CC Kit for assaying MDR1 gene expression level

FT SOURCE

1. 120

FT SOURCE

/organism='Artificial Sequence'.
 BD160756 120
Kit for assaying MDR1 gene e
BD160756
BD160756.1 GI:27866514
JP 2002153274-A/4.
synthetic construct
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Kit for a
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1 (bases 1 to 120)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       multiple drug resistance
Patent: WO 0234291-A 29 02-MAY-2002;
THE BRIGHAM AND WOMEN'S HOSPITAL, INC.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                          Similarity
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.larity 100.0%; !
Conservative 0;
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                                                                                                                                                                                                                                                                                        /organism='Artificial
                                                                                                                                                                                                                                       /mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                               organism="synthetic construct"
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/mol_type="unassigned DN
/db_xref="taxon:9606"
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                                                                                                                                                                          Score 25; DB
Pred. No. 0.0
D; Mismatches
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RESULT 6
CQ148922
LOCUS
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AUTHORS
TITLE
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CQ110210
LOCUS
DEFINITION
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AUTHORS
TITLE
JOURNAL
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CQ148922
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llarity 100.0%;
Conservative 0
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DNA
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ISM synthetic construct
other sequences; artificial sequences.

CE 1 (bases 1 to 120)

I (bases 1 to 120)

Kit for assaying MDR1 gene expression level
AL For assaying MDR1 gene expression level
AL RAINOS LAB INC

OS Artificial Sequence
PN JP 2002153274-A/4

PD 28-MAY-2002

FT 17-NOV-2000 JP 2000351224
PF 17-NOV-2000 JP 2000351224
PF TAKESHI UUIIE, TSUKASA HAYASHI, MASAKO KAWASE, FUMIAKI UEJI PC

C12NN15/99,C12N15/00

CC Kit for assaying MDR1 gene expression level
FT Key Location/Qualifiers
FT Source /organism='Artificial Sequence'.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               172 bp DNA Sequence 19069 from Patent WO0157272. CQ110210 CQ110210.1 GI:41079573
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Penn, S.G., Hanzel, D.K., Chen, W. and Rank, D.R. Human genome-derived single exon nucleic acid analysis of gene expression in human placenta Patent: WO 0157272-A 19069 09-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
                                                                                                                                                                                                                                                                                                                                                                  /organism="Homo sapiens"
/mol type="unassigned DNA"
/mol type="unassigned DNA"
/db xref="teaxon:9606"
/noTe="MAP TO AC005068.1-EXPRESSED IN PLACENTA, SIGNAL
/noTe="MAP TO AC005068.1-EXPLUE 3.00e-27-NT HIT:
AF016535.1, EVALUE 2.00e-92~EST_HUMAN HIT: AW847648.1,
EVALUE 2.30e+00"
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/mol_type="genomic DNA"
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Pred. No. 0.055;
Mismatches
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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                     HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR ANALYSIS OF GENE EXPRESSION IN HUMAN ADULT LIVER-(130) PB 0004 WO 3<150> US 60/180,312<151> 04 February 2000 (04.02.00)<150> US 60/180,312<151> 04 February 2000 (04.02.00)<150> US 60/207,45<4151> 26 May 2000 (26.05.00)<150> US 09/632,36<4151> 03 August 2000 (03.08.00)<150> GB 24263.6<151> 03 October 2000 (03.10.00)<150> US 60/236,359<4151> 27 September 2000 (27.09.00)<150> US 60/236,687<4151> 21 September 2000 (21.09.00)<150> US 09/608,408<4151> 30 June 2000 (30.06.00)<170> Wolecular Dynamics Sequence Listing Engine Learning Engine Lear
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Human genome-derived single exon nucleic acid probes useful
analysis of gene expression in human bone marrow
patent: WO 0157276-A 18944 09-AUG-2001;
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CQ148922
ch 100.0%;
l Similarity 100.0%;
25; Conservative 0
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                           /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
/note="MAP TO ACOOS068.1.~EXPRESSED IN ADULT LIVER, SIGNAL
= 0.73-NT HIT: AF016535.1, EVALUE 2.00e-92~SWISSPROT HIT:
P08183, EVALUE 3.00e-27-EST_HUMAN HIT: AW847648.1, EVALUE
2.30e+00"
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/mol type="unasaigned DNA"
/db xref="taxon:9606"
/db xref="taxon:9606"
/note="MAP TO AC005068.1~EXPRESSED IN BONE MARROW, SIGNAL
- 0.33~SWISSPROT HIT: P08183, EVALUE 3.00e-27~NT HIT:
aF016535.1, EVALUE 2.00e-92~EST_HUMAN HIT: AW847648.1,
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Pred. No.
                               Score 25; DB 6;
Pred. No. 0.055;
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Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 25; Conservative 0;
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CQ307508
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                Penn, S.G., Hanzel, D.K., Chen, W. and Rank, D.R. Human genome-derived single exon nucleic acid analysis of gene expression in human brain Patent: WO 0157275-A 18468 09-AUG-2001;
                                                                                                                                                                                                                                                                                                                                           Homo sapiens
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/db_xref="taxon:9606"
/note="MAP TO AC005068.1-EXPRESSED IN LUNG, SIGNAL = 0.65-SWISSPROT HIT: P08183, EVALUE 3.00e-27-NT HIT: AF016535.1, EVALUE 2.00e-92~EST_HUMAN HIT: AW847648.1, EVALUE 2.30e+00"
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/db_xref="taxon:9606"
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                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 201)
Nelson,R.M., Marnellos,G., Kammerer,S., Hoyal,C.R., Shi,M.M., Cantor,C.R. and Braun,A.
Large-Scale Validation of Single Nucleotide Polymorphisms in Gene
                                                                                                                                                                                                                         201 bp sqrm208304 Human DNA (Sequenom) tagged site.
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             Genome Res.
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BV202142
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3595 John Hopkins Court, San Diego,
Tel: 18582029018
                                                                                                                                                            Homo sapiens (human)
                                                                                                                                                                                            BV202142.1 GI:48171156
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sqnm208303 Human DNA (Sequenom)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: abraun@sequenom.com
Primer A: No primer sequence submitted
Primer B: No primer sequence submittéd
STS size: 201.
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Cantor,C.R. and Braun,A.
Large-Scale Validation of Single Nucleotide Polymorphisms in Gene
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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vative 0; Mismatches
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1 (bases 1 to 209)

Gekeler, V., Weger, S. and Probst, H.

mdr1/P-glycoprotein gene segments analyzed from various human leukemic cell lines exhibiting different multidrug resistance
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M37724.1 GI:183537
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3595 John Hopkins Court, San Diego,
Tel: 18582029018
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Primer A: No primer sequence submitted
Primer B: No primer sequence submitted
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complement(1. .19)
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<1...>201
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                                                                                                                                                                                     gene="PGY1"
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type="T-lymphoblast"
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CQ149621/c
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CQ308221/c
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Sequence 19326 from Patent W00186003.
CQ308221
CQ308221.1 GI:41268798
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Human genome-derived single exon nucleic acid probes useful for
analysis of gene expression in human bone marrow
Patent: WO 0157276-A 19643 09-AUG-2001;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CQ149621 219 bp DNA Sequence 19643 from Patent WO0157276. CQ149621 GI:41156971
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                                                                                     Human genome-derived single exon nucleic acid snalysis of gene expression in human lung patent: WO 0186003-A 19326 15-NOV-2001;
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25; Conservative
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Search completed: February Job time: 483.178 secs

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CQ345026/c
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Sequence 19120 from Patent WO0157275.
CQ345026 CQ345026.1 GI:41294097
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                                                                        1 AGCTTCCAACCACGTGTAAATCCTA 25
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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P08183, EVALUE 2.00e-27"
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520.498 Million cell updates/sec
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US-08-784-649A-5
US-08-181-471-2
US-08-181-471-1
US-09-136-167-1
US-09-37-233-1
S206352-3
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US-09-397-610-6
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US-09-672-725C-24
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RESULT 1
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;Patent No. 52063
; APPLICANT: F
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; TITLE OF INVENTION: COMPOSITIONS FOR CLONES CONTAINING DNA
; SEQUENCES ASSOCIATED WITH MULTIDRUG RESISTANCE IN HUMAN CELLS
; NUMBER OF SEQUENCES: 4
; CURRENT APPLICATION DATA:
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TITLE OF INVENTION: COMPOSITIONS FOR CLONES CONTAINING DNA
SEQUENCES ASSOCIATED WITH MULTIDRUG RESISTANCE IN HUMAN CELLS
NUMBER OF SEQUENCES: 4
CURRENT APPLICATION DATA:
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APPLICANT: Roninson,
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LENGTH: 807
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Matches 25; Conserv
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FILING DATE: 24-SEP-1990
PRIOR APPLICATION DATA:
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PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 892,575
PILING DATE: 01-AUG-1986
APPLICATION NUMBER: 845,610
PILING DATE: 28-MAR-1986
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FILING DATE: 28-MAR-1986
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                  807
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                              Igor B.;Pastan Ira H.;Gottesman,
                                                                                                                                                                              US/07/622,836
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US-09-949-016-71318
US-09-949-016-179555
US-09-949-016-179556
US-09-949-016-179564
US-09-949-016-17437
US-09-949-016-15879
US-09-949-016-15921
US-09-949-016-17533
US-09-949-016-30006
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Pred. No. 0.0074
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664, APP
12437, A
11935, A
16935, A
16935, A
17533, A
17533, A
12371, A
1237
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71318,
86805,
179555,
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Query Match

100.0%;

Score 25;

DB 6;

Length 807;

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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-762-195-1
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                                                                                                                                                                             SEQ ID NO 1
LENGTH: 4186
TYPE: DNA
    Best Loc
Matches
  Ouery Match 100.0%; Score 25; DB 4; Length 4186; Best Local Similarity 100.0%; Pred. No. 0.0097; Matches 25; Conservative 0; Mismatches 0; Indels 0.
                                                                                                                                                                                                                                                                                         CURENT APPLICATION NUMBER: US/09/672,810
CURRENT FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US 60/156,921
PRIOR FILING DATE: 1999-09-79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1, Application US/09672810 Patent No. 6617450
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APPLICANT: STEINMEL-CRESPI, DOI
APPLICANT: CRESPI, CHARLES L.
APPLICANT: CRESPI, CHARLES L.
TITLE OF INVENTION: P-GLYCOPRI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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APPLICANT: Stremmel, Wolfgang
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                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US PRIOR FILING DATE: 1999-10-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: PCT/EP99702426
PRIOR FILING DATE: 1999-08-06
PRIOR APPLICATION NUMBER: 198 35 526 2 DE
PRIOR PILING DATE: 1998-08-06
PRIOR PPLICATION NUMBER: 198 57 570.8 DE
PRIOR FILING DATE: 1998-12-15
PRIOR FILING DATE: 1998-12-15
PRIOR FILING DATE: 1998-12-15
PRIOR FILING DATE: 1998-12-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Phosphatidylcholine as Medication with TITLE OF INVENTION: Protective Effect on Large Intestinal Mucosa FILE REFERENCE: 34691/208520 CURRENT APPLICATION NUMBER: US/09/762,195 CURRENT FILING DATE: 2001-02-05
                                                                                              NAME/KEY: CDS
LOCATION: (100)...(3940)
                                                                                                                                                        ORGANISM: Macaca
                                                                                                                                        FEATURE:
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5. 6677319
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STOCKER, PENNY J.
STOCKER, PENNY J.
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                                                                                                                                                                                                                                                                                                                                                                                                            P-GLYCOPROTEINS AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Pred. No. 0.0074; tive 0; Mismatches 0;
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0.0097;
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; FEATURE:
; NAME/KEY: CDS
; LOCATION: (100
US-09-672-810-3
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US-08-784-649A-1/c
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CURRENT APPLICATION NUMBER: US/09/672,810
CURRENT FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US 60/156,921
PRIOR FILING DATE: 1999-09-28
PRIOR APPLICATION NUMBER: US 60/158,818
PRIOR FILING DATE: 1999-10-12
NUMBER OF SEQ ID NOS: 18
SOFTWARE: FastSEQ for Windows Version 3.0
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US-09-672-810-3/c
SOFTWARE: PATENTIN Release #1.0, Versio CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/784,649A
FILING DATE:
CLASSIFICATION: 435
ATTORNEY_AGENT INFORMATION:
NAME: Sherwood, Pamela J
REGISTRATION NUMBER: Reg No. 5830697 36
REFERENCE_DOCKET NUMBER: 06037/007001
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: STOCKER, PENNY J.
APPLICANT: STEIMEL-CRESPI, DOROTHY T.
APPLICANT: CRESPI, CHARLES L.
TITLE OF INVENTION: P-GLYCOPROTEINS AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA ORGANISM: Macaca fascicularis
                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
STREET: 2200 Sand Hill Road
                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Sikic, Brancher APPLICANT: Chen, Gang
APPLICANT: Chen, Gang
APPLICANT: Chen, Gang
APPLICANT: MODULATION TO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ENGTH: 4195
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                                                                                                                                                                                                                                                                                                                                CITY: Menlo Park
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 749
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5. 5830697
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                                                                                                                                                                                                                                                                            94025
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Similarity 100.0%;
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                                                                                                                                                                           Version #1.25
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TELEFAX: 415-854-0875
INFORMATION FOR SEQ ID NO:

SEQUENCE CHARACTERISTICS

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RESULT 7
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RESULT 8
US-08-181-471-2/c
; Sequence 2, Application US/08181471
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                                                                                                                                                                                                                                         US-08-784-649A-5
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LENGTH: 4264 bass
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Patent No. 5830697
GENERAL INFORMATION:
APPLICANT: Sikic, Branimir I
APPLICANT: Chen, Gang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 100.0%; Score 25; DB 2; Best Local Similarity 100.0%; Pred. No. 0.0098; Matches 25; Conservative 0; Mismatches C
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Best Local Similarity
Matches 25; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/784,649A
FILING DATE:
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Sherwood, Pamela J
REGISTRATION NUMBER: Reg.No. 5830697 36,677
REFERENCE/DOCKET NUMBER: 06037/007001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-322-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
STREET: 2200 Sand Hill Road
CITY: Menlo Park
                                                                                                                                                                                                                                                           MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TLE OF INVENTION: F-GLYCOPROTEIN MUTANT RESISTANT TO TILE OF INVENTION: CYCLOSPORIN MODULATION
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TOPOLOGY: lir
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                                                                                                AGCTTCCAACCACGTGTAAATCCTA 755
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100.0%; Pred. No. 0.0098;
tive 0; Mismatches 0
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RESULT 9
US-09-023-655-1167/c
; Sequence 1167, Application US/09023655
; Patent No. 6607879
; Patent No. 6607870N:
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 619-792-8477
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Li, Lingna
APPLICANT: Lishko, Val
TITLE OF INVENTION: ME
TITLE OF INVENTION: CO
NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: Patentin Rel-Base #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Thomas Fitting
STREET: 12526 High Bluff Drive, Suite 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 4646 base pair
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: AN TELECOMMUNICATION INFORMATION: TELEPHONE: 619-792-3680
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 02-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Fitting, Thomas
REGISTRATION NUMBER: 34,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                             APPLICANT: Cocks, Benjamin G. APPLICANT: Susan G. Stuart APPLICANT: Jeffrey J. Seilhamer TITLE OF INVENTION: COMPOSITION TITLE OF INVENTION: EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ANTI-SENSE:
                                                               CORRESPONDENCE ADDRESS:
                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/0 FILING DATE: 13-JAN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: San Diego
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
CITY: PALO ALTO
STATE: CALIFORNI
                                STREET:
                                              ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
                                                                                                                                                                                                                                                                                              1065 AGCTTCCAACCACGTGTAAATCCTA 1041
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 CALIFORNIA
                            E: INCYTE PHARMACEUTICALS,
3174 PORTER DRIVE
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425..4267
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METHOD FOR DELIVERING BENEFICIAL
COMPOSITIONS TO HAIR FOLLICLES
                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 25; 100.0%; Pred. No.
                                                                                 1508
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                                                                                                                THE DETECTION
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                                                                                                                  BLOOD CELL GENE
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COMPUTER READABLE FORM:

94304

MEDIUM TYPE:

Floppy disk

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; TOPOLOGY: linear ; IMMEDIATE SOURCE; ; LIBRARY: GENBANK ; CLONE: 9187468 US-09-023-655-1167
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (650) 845-4166
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/332,444
FILING DATE: 31-OCT-1994
                                                                                                                                                                                                                                  STREET: 6 LL.
STREET: New Jersey
                                                                         SOFTWARE: DW4.V2
SOFTWARE: DW4.V2
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/583,276
FILING DATE: 05-JAN-1996
                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Zeller, Karen J.
REGISTRATION NUMBER: 37,071
REFERENCE/DOCKET NUMBER: PA-
TELECOMMUNICATION INFORMATION:
TELECPHONE: (650) 855-0555
                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Carella,
                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
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                                                                                                                                  COMPUTER: 18M PS/2
OPERATING SYSTEM: PC-DOS
SOFTWAPF. ...
                                                                                                                                                                                                                     COUNTRY: US
ZIP: 07068
                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                               TLE OF INVENTION:
TLE OF INVENTION:
                                                                                                                                                                                                                                                                                                           ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS 6.2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nucleic acid
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                                                                                                                                                                                                                                                                                         6 Becker Farm Road
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Nienhuis, Arthur
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                                                                                                                                                                                                                                                                                                           Carella, Byrne, Bain, Gilfillan,
Cecchi & Stewart
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MULTIDRUG RESISTANCE GENES AND IMPROVED
SELECTION OF CELLS TRANSDUCED WITH SUCH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kevin T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            us/09/023,655
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US-08-752-447-1
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US-08-752-447-1/c
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                                                                                                                                                                                                  TELEFAX: 312-913-9808
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 4669 base pair
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 07/8 FILING DATE: 22-MAY-1992 INFORMATION FOR SEQ ID NO: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                           FEATURE:
                                                                                                                                                                                                                                                               TELEPHONE: 312-913-0001
                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: NO. 5994088nan, Kevin
REGISTRATION NUMBER: 35,303
REFERENCE/DOCKET NUMBER: 95
                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE
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LENGTH: 4669 bases
                                                                                                                                             MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
            NAME/KEY:
                                                                                                                                                             STRANDEDNESS: si
TOPOLOGY: linear
                                                        LOCATION:
                                                                                                                NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Chicago
STATE: Illinois
                                                                                                                                                                                        TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/0 FILING DATE: 15-NOV-1996 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
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STRANDEDNESS: singular
                                                                                                    OCATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DESCRIPTION:
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                                                                                                                                                                                                      4669 base pairs
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4265..4669
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425..4264
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Pred. No. 0.0099;
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Query Match

100.0%;

Score 25;

DB 2;

Length 4669;

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RESULT 12
US-09-316-167-1/c
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RESULT 13
US-09-397-233-1/c
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                                                                                                                                                                                             US-09-316-167-1
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                                                                                                                           Matches
                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION UNMBER: 08/752,44
PILING DATE: 15-NOV-1996
ATTORNEY/AGENT INFORMATION:
NAME: No. 6365357nan, Kevin E
REGISTRATION NUMBER: 35,303
REFERENCE/DOCKET NUMBER: 95,1
TELECOMMUNICATION INFORMATION:
TELECHONE: 312-913-0001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Mechetner, Eugene
APPLICANT: Roninson, Igor B
TITLE OF INVENTION: Methods and Reagents for Preparing
TITLE OF INVENTION: Using Immunoligaal Agents Specific
NUMBER OF SEQUENCES: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Rel-Base #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff Ltd
STREET: 300 South Wacker Drive, Seventh Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                        FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 300
STREET: Chicago
                                                                                                                                                                                                                                                                                                                                                   OLECULE TYPE:
                                                                                                                                                                                                          NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: FILING DATE:
                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: USA
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                                                                                                                                                                                                                                                                          NAME/KEY:
                                                                                                                                                                                                                                                                                                                         NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 4669 base pairs
                                                                                                                                                                                                                                                                                                        LOCATION:
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                                                                                                                                            Similarity
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                                                               AGCTTCCAACCACGTGTAAATCCTA 1041
                                                                                     AGCTTCCAACCACGTGTAAATCCTA 25
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425..4264
                                                                                                                             Conservative
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1..424
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4265..4669
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                                                                                                                          100.0%; Score 25; DB 3; Length 4669; 100.0%; Pred. No. 0.0099; tive 0; Mismatches 0; Indels
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for P-glycoprotein
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LOCATION: 4265..4669
SEQUENCE DESCRIPTION: SEQ ID NO: 1:
US-09-397-233-1
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5206352-3/c
;Patent No. 5206352
; APPLICANT: Roninson, Igor B.;Pastan Ira H.;Gottesman,
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                                                                                                                        ;Michael M.
TITLE OF INVENTION: COMPOSITIONS FOR CLONES CONTAINING DNA;SEQUENCES ASSOCIATED WITH MULTIDRUG RESISTANCE IN HUMAN CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                Matches
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 892,575
FILING DATE: 01-AUG-1986
                                                                                        NUMBER OF SEQUENCES: 4
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/09/397,233

FILING DATE: 16-Sep-1999

CLASSIFICATION: «Unknown»

ATTORNEY/ACENT INFORMATION:

NAME: No. 6630327nan, Kevin E

REGISTRATION NUMBER: 35,303

REFERENCE/DOCKET NUMBER: 95,1121-C

TELECOMMUNICATION INFORMATION:

TELEPHONE: 312-913-0001
                                                      APPLICATION NUMBER: US/
FILING DATE: 24-SEP-1990
                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTMARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Roninson, Igor B
TITLE OF INVENTION: Methods and Reagents for Preparing and
Using Immunological Agents Specific fo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                           1065 AGCTTCCAACCACGTGTAAATCCTA 1041
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                                                                                                 1 AGCTTCCAACCACGTGTAAATCCTA 25
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LOCATION:
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STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY:
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1..424
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425..4264
                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 25; DB 4; 100.0%; Pred. No. 0.0099
                                                                           US/07/622,836
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RESULT 15
5206352-3/c
;Patent NO. 5206352
; ARPLICANT: RONINBON, IGOT B.;Pastan Ira H.;Gottesman,
;Michael M.
TITLE OF INVENTION: COMPOSITIONS FOR CLONES CONTAINING DNA
;SEQUENCES ASSOCIATED WITH MULTIDRUG RESISTANCE IN HUMAN CELLS
NUMBER OF SEQUENCES: 4
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/622,836
FILING DATE: 24-SEP-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 992,575
FILING DATE: 01-AUG-1986
; APPLICATION NUMBER: 845,610
; FILING DATE: 28-MAR-1986
; SEQ ID NO:3:
LENGTH: 4669
520632-1
Search completed: Pebruary 9, 2005, 17:11:17 Job time: 79.592 secs
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FILING DATE: 28-MAR-1986
;SEQ ID NO:3:
LENGTH: 4669
                                                                                                                                                           Query Match 100.0%; Score 25; DB 6; Length 4669; Best Local Similarity 100.0%; Pred. No. 0.0099; Matches 25; Conservative 0; Mismatches 0; Indels 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 100.0%; Score 25; DB 6; Length 4669; Best Local Similarity 100.0%; Pred. No. 0.0099; Matches 25; Conservative 0; Mismatches 0; Indels
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Title:
Perfect score:
Sequence:
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Scoring table:
Published Applications NA:*

1: /cgm2_6/prodata/1/pubpna/US07_PUBCOMB.seq:*

2: /cgm2_6/prodata/1/pubpna/US06_NEW_PUB.seq:*

3: /cgm2_6/prodata/1/pubpna/US06_NEW_PUB.seq:*

4: /cgm2_6/prodata/1/pubpna/US07_NEW_PUB.seq:*

5: /cgm2_6/prodata/1/pubpna/US07_NEW_PUB.seq:*

6: /cgm2_6/prodata/1/pubpna/US08_NEW_PUB.seq:*

7: /cgm2_6/prodata/1/pubpna/US08_NEW_PUB.seq:*

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14: /cgm2_6/prodata/1/pubpna/US09E_PUBCOMB.seq:*

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20: /cgm2_6/prodata/1/pubpna/US10E_PUB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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W N | , , , | Result |
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| 25
25 | 25 | 2 2 5 5 | 255 | 25 5 | 25 | Score |
| 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | Query
Match |
| 3860
4186 | 3860 | 3840 | . 473 | 219 | 172 | Query
Match Length DB |
| 17 | 9 <u>1</u> 5 | 18 | · • • | 9 5 | ۰ | BB |
| US-09-866-866A-3
US-10-619-359A-1 | US-10-101-433A-1
US-09-866-866A-1 | US-09-805-020-31
US-10-384-339C-30 | US-09-864-761-10820
US-09-864-761-11142 | US-10-101-510-284
US-09-864-761-27770 | US-09-864-761-27462 | ID |
| Sequence 3, Appli
Sequence 1, Appli | Sequence 1, Appli | sequence 31, Appl
Sequence 30, Appl | Sequence 10020, A
Sequence 11142, A | Sequence 27770, A | Sequence 27462, A | Description |

ALIGNMENTS

| ACID PROBES USEFUL FOR | PRIOR APPLICATION NUMBER: PCT/USO1/00667 PRIOR FILING DATE: 2001-01-30 PRIOR RPLICATION NUMBER: PCT/USO1/00664 PRIOR PLICATION NUMBER: PCT/USO1/00669 PRIOR APPLICATION NUMBER: PCT/USO1/00669 PRIOR PILING DATE: 2001-01-30 PRIOR APPLICATION NUMBER: PCT/USO1/00665 PRIOR APPLICATION NUMBER: PCT/USO1/00665 PRIOR FILING DATE: 2001-01-30 |
APPLICANT: Penn, Sharron G. APPLICANT: Penn, David R. APPLICANT: Hanzel, David K. APPLICANT: Chen, Wenaheng TITLE OF INVENTION: GENO EXPRESSION ANALYSIS BY MICROARRAY FILE AFFERBYCE: Acomica-X-1 CURRENT APPLICATION NUMBER: US/09/864,761 CURRENT FILING DATE: 2001-05-23 PRIOR APPLICATION NUMBER: US 60/180,312 PRIOR APPLICATION NUMBER: US 60/207,456 | ESULT 1
S-09-864-761-27462
Sequence 27462, Application US/09864761
Patent No. US20020048763A1
CENTERAL TUROPMATION: |
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APPLICATION NUMBER: PCT/US01/00668

FOR

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US-10-101-510-284
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Matches
                                                                                      Query Match
Best Local Similarity
Matches 25; Conserv
                                                                                                                                                                                                                                              SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 284
LENGTH: 209
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: WAN, JACKSON
APPLICANT: WANG, YIXIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 284, Appropriate Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: EXPRESSION PROFILES AND METHODS OF USE FILE REFERENCE: 15117.0012
CURRENT APPLICATION NUMBER: US/10/101,510
CURRENT FILING DATE: 2002-03-20
PRIOR APPLICATION NUMBER: 60/276,947
PRIOR FILING DATE: 2001-03-20
                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: MAP TO AC005068.1

OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 0.64

OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 0.65

OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 0.53

OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 0.73

OTHER INFORMATION: EXPRESSED IN PLACEBURA, SIGNAL = 0.64

OTHER INFORMATION: SWISSPROT HIT: P08183, EVALUE 3.00e-27

OTHER INFORMATION: NT HIT: AF016535.1, EVALUE 2.00e-92

OTHER INFORMATION: EST_HUMAN HIT: AW847648.1, EVALUE 2.30e+00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ENGTH: 172
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OR APPLICATION NUMBER: PCT/USO1/00663
OR FILING DATE: 2001-01-30
OR FILING DATE: 2001-01-30
OR PILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/USO1/00661
OR APPLICATION NUMBER: PCT/USO1/00661
OR APPLICATION NUMBER: PCT/USO1/00670
OR APPLICATION NUMBER: PCT/USO1/00670
OR APPLICATION NUMBER: PCT/USO1/00670
OR FILING DATE: 2001-01-30
OR FILING DATE: 2000-09-21
OR APPLICATION NUMBER: US 09/608,408
OR APPLICATION NUMBER: US 09/774,203
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  204
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AGCTTCCAACCACGTGTAAATCCTA 180
                                       AGCTTCCAACCACGTGTAAATCCTA 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Application US/10101510 to. US20030148295A1
                                                                                    100.0%; Score 25; Dilarity 100.0%; Pred. No. 0.0 Conservative 0; Mismatches
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                                                                                                         DB 15; Length 209; 0.041;
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; OTHER INFORMATION: MAP TO AC002457.1
; OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 0.42
; OTHER INFORMATION: EXPRESSED IN BOME MARROW, SIGNAL = 0.57
; OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 0.57
; OTHER INFORMATION: NT HIT: M29428.1, EVALUE 1.00e-120
; OTHER INFORMATION: SWISSPROT HIT: P08183, EVALUE 2.00e
US-09-864-761-27770
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CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 90/632,366
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 90/632,366
PRIOR FILING DATE: 2000-08-03
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Annomax Sequence
SEQ ID NO 27770
LENGTH: 219
                                                                                                                                 Matches
                                                                                                                                                                              Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR FILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR FILING DATE: 2001-01-29
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APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
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OR APPLICATION NUMBER: PCT/US01/00664
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00669
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00665
OR APPLICATION NUMBER: PCT/US01/00665
OR FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00670
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00662
FILING DATE: 2001-01-30
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                                                                                                                                                                              Similarity
AGCTTCCAACCACGTGTAAATCCTA 25
                                                                                                                       100.0%;
larity 100.0%;
Conservative 0
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Pred. No. 0.041;
; Mismatches
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RESULT 4
US-09-864-761-10820
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Best Local S
Matches 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
SEQ ID NO 10820
LENGTH: 463
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CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
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APPLICANT: Hanzel, David K.
APPLICANT: Chen, Mensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
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PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
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                                                                                                                                                                                                                       OTHER INFORMATION: MAP TO AC005068.1

OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 0.64

OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 0.65

OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 0.53

OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 0.73

OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 0.64
                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
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OR FILING DATE: 2001-01-30

DR APPLICATION NUMBER: PCT/US01/00664

DR FILING DATE: 2001-01-30

DR APPLICATION NUMBER: PCT/US01/00665

DR APPLICATION NUMBER: PCT/US01/00665

DR APPLICATION NUMBER: PCT/US01/00665
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APPLICATION NUMBER: PCT/US01/00661
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00670
FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/USO1/00668
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/USO1/00663
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/USO1/00662
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APPLICATION NUMBER: PCT/US01/00666
FILING DATE: 2001-01-30
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APPLICATION NUMBER: GB 24263.6
FILING DATE: 2000-10-04
APPLICATION NUMBER: US 60/236,359
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APPLICATION NUMBER: US 60/207,456
FILING DATE: 2000-05-26
APPLICATION NUMBER: US 09/632,366
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25; Conserv
     AGCTTCCAACCACGTGTAAATCCTA 364
                                         AGCTTCCAACCACGTGTAAATCCTA 25
                                                                                             100.0%;
ilarity 100.0%;
Conservative 0
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Pred. No. 0.047;
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RESULT 5
US-09-864-761-11142/c
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PRIOR FILING DATE: 2000-10-04
PRIOR PRILING DATE: 2000-10-04
PRIOR PRILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR PRILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
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                                                                                                                                                                       OTHER INFORMATION: MAP TO AC002457.1
OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 0.42
OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL:
OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 0.57
US-09-864-761-11142
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LENGTH: 473
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Best Local
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PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 2000-08-03
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CURRENT FILING DATE: 2001-05-23
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NUMBER OF SEQ ID NOS: 49117
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PRIOR APPLICATION NUMBER: US 60/234,687
                                                                                                                                                                                                                                                                                                   LENGTH: 473
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                     FEATURE:
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00670
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                           AGCTTCCAACCACGTGTAAATCCTA 25
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Hanzel, David K.
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ilarity 100.0%;
Conservative (
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: 2001-01-29
IOS: 4017
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                                                                                                           Score 25;
Pred. No.
                                                                                       Mismatches
                                                                                                              DB 9;
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                                                                                                                                  Length 473;
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RESULT 8
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; PATENT DOCUMENT NUMBER: AF016535
US-10-384-339C-30
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CURRENT APPLICATION NUMBER: US/10/384,339C

CURRENT FILING DATE: 2003-03-07

PRIOR APPLICATION NUMBER: PCT/EP02/00152

PRIOR FILING DATE: 2002-01-09

PRIOR FILING DATE: 2001-01-09

PRIOR FILING DATE: 2001-10-26

PRIOR APPLICATION NUMBER: DE 10155280.7

PRIOR FILING DATE: 2001-10-26

PRIOR FILING DATE: 2001-11-26

PRIOR APPLICATION NUMBER: DE 10158411.3

PRIOR APPLICATION NUMBER: DE 10158411.4

PRIOR APPLICATION NUMBER: DE 10160151.4

PRIOR APPLICATION NUMBER: DE 10160151.4
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US-10-384-339C-30/c
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US-09-805-020-31/c
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publication No. US20020086384A1
GENERAL INFORMATION:
APPLICANT: LEVINE, Zurit
TITLE OF INVENTION: SPLICE VARIANTS OF ONCOGENES
FILE REFERENCE: 2786-0168P
CURRENT APPLICATION UNMERR: US/09/805,020
CURRENT FILING DATE: 2001-03-13
NUMBER OF SEQ ID NOS: 72
                                                                                                                                                        Matches
                                                                                                                                                                                                 Query Match
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                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Homo sapiens
PUBLICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 2001-12-07
NUMBER OF SEQ ID NOS: 173
                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Ver.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Kreutzer, Roland
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR INHIBITING EXPRESSION OF A TARGET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc_feature
LOCATION: (1)..(2307)
OTHER INFORMATION: any n = a,c,g,t any unknown
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TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                          ENGTH: 3840
                                                                                                                                                                          Local
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                                                                                          1 AGCTTCCAACCACGTGTAAATCCTA 25
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                                                                                                                                                                             Similarity
                                                              AGCTTCCAACCACGTGTAAATCCTA 617
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100.0%; Pred. No. 0.
ative 0; Mismatcher
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RESULT 10 US-09-866-866A-3/c

GENERAL INFORMATION

APPLICANT: Sorrentino, Bri APPLICANT: Schuetz, John

Brian

Sequence 3, Application US/09866866A Patent No. US20020102244A1

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                                                                                                                                                                      ; LENGTH: 3860
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-866A-1
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                                                                                     Matches
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                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
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Best Local
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                                                                                                                                                                                                                                                                                                                APPLICANT: Schuetz, John
TITLE OF INVENTION: A Method of Identifying and/or Isolating Stem
FILE REFERENCE: 1340-1-021CIP2
CURRENT APPLICATION NUMBER: US/09/866,866A
CURRENT FILING DATE: 2001-08-30
PRIOR APPLICATION NUMBER: 09/584,586
PRIOR FILING DATE: 2000-05-31
PRIOR APPLICATION NUMBER: PCT/US99/11825
PRIOR APPLICATION NUMBER: PCT/US99/11825
PRIOR FILING DATE: 1999-05-27
PRIOR PILING DATE: 1999-05-27
PRIOR FILING DATE: 1998-05-28
PRIOR FILING DATE: 1998-05-28
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APPLICANT: Crespi, Charles
TITLE OF INVENTION: P-GLYCOPROTEINS AND USES THEREOF
FILE REFERENCE: G00307/70019
CURRENT APPLICATION NUMBER: US/10/101,433A
CURRENT FILING DATE: 2002-03-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Sorrentino, Brian APPLICANT: Schuetz, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 60/277,095
PRIOR FILING DATE: 2001-03-19
NUMBER OF SEQ ID NOS: 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Macaca mulatta FEATURE:
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LOCATION: (1)..(3852)
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641 AGCTTCCAACCACGTGTAAATCCTA 617
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                           AGCTTCCAACCACGTGTAAATCCTA 25
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                                                                                   Conservative
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                                                                                                     100.0%; Score 25; DB 9; 100.0%; Pred. No. 0.068;
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100.0%; Pred. No. 0.068;
tive 0; Mismatches
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                                                                                   Mismatches
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APPLICANT: CRESPI, CHARLES L.
TITULE OF INVENTION: P-GLYCOPROTEINS AND USES THEREOF
FILE REFERENCES: G00307.70020. US
CURRENT APPLICATION NUMBER: US/10/619,359A
CURRENT FILING DATE: 2003-07-14
PRIOR APPLICATION NUMBER: US 60/156,921
PRIOR FILING DATE: 1999-09-28
PRIOR PRIOR APPLICATION NUMBER: US 60/158,818
PRIOR APPLICATION NUMBER: US 60/158,818
PRIOR APPLICATION NUMBER: US 69/672,810
PRIOR FILING DATE: 1999-10-12
PRIOR FILING DATE: 2000-9-28
                                                        RESULT 12
US-10-619-359A-3/c
US-10-619-359A-3/c
J. Sequence 3, Application US/10619359A
J. Publication No. US20040077000A1
J. Publication No. US20040077000A1
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CURRENT FILING DATE: 2001-08-30
PRIOR APPLICATION NUMBER: 09/584,586
PRIOR FILING DATE: 2000-05-31
PRIOR APPLICATION NUMBER: PCT/US99/11825
PRIOR FILING DATE: 1999-05-27
PRIOR APPLICATION NUMBER: 60/086,988
PRIOR FILING DATE: 1998-05-28
NUMBER OF SEQ ID NOS: 27
SOFTWARE: Patentin version 3.0
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                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: STOCKER, PENNY J.
APPLICANT: STEIMEL-CRESPI, DOROTHY T.
APPLICANT: CRESPI, CHARLES L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: A Method of Identifying and/or Isolating Stem Cells FILE REFERENCE: 1340-1-021CIP2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 18 SOFTWARE: FastSEQ for Windows Version 3.0
                  APPLICANT:
  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 3860
TYPE: DNA
ORGANISM: homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 4186
TYPE: DNA
ORGANISM: Macaca fascicularis
FEATURE:
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LOCATION: (100)...(3940)
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STOCKER, PENNY J.
STEIMEL-CRESPI, DO
CRESPI, CHARLES L.
                                                                                                                                                                                                     AGCTTCCAACCACGTGTAAATCCTA 716
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llarity 100.0%; Pred. No. 0.
Conservative 0; Mismatches
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ilarity 100.0%;
Conservative 0
                    DOROTHY T
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Pred. No. 0.069;
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Query Match
Best Local Similarity
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US-10-072-621-2/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: LEVINE, ZURIL
APPLICANT: LEVINE, SPLICE VARIANTS OF ONCOGENES
FILE REFERENCE: 2786-0168P
CURRENT APPLICATION NUMBER: US/09/805,020
CURRENT FILING DATE: 7201-03-13
NUMBER OF SEQ ID NOS: 72
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 18
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 3
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                                                                                                                             GENERAL INFORMATION:
                                                                                                                                              Sequence 2, Application US/10072621 Publication No. US20020169137A1
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CURRENT FILING DATE: 2003-07-14
PRIOR APPLICATION NUMBER: US 60/156,921
PRIOR FILING DATE: 1999-09-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: US 60/158,818
PRIOR FILING DATE: 1999-10-12
PRIOR APPLICATION NUMBER: US 09/672,810
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                         APPLICANT: Reiner, Peter B.
APPLICANT: Connop, Bruce P.
APPLICANT: Connop, Bruce P.
APPLICANT: Pollard, Michalle
TITLE OF INVENTION: REGULATION OF AMYLOID PRECURSOR PROTEIN EXPRESSION
TITLE OF INVENTION: BY MODIFICATION OF ABC TRANSPORTER EXPRESSION OR ACTIVITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA ORGANISM: Macaca fascicularis FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: CDS
LOCATION: (100)...(3949)
CURRENT APPLICATION NUMBER: US/10/072,621
                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: misc_feature LOCATION: (1)..(4533)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
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                                                                                                                                                                                                                                                                1065 AGCTTCCAACCACGTGTAAATCCTA 1041
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Pred. No.
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APPLICANT: Karen LU

APPLICANT: Karen LU

APPLICANT: Karen ZHAT

APPLICANT: Xumei ZHAT

APPLICANT: Karen ZHAT

ITILE OF INVENTION: Nucleic Acid Molecules and Proteins For The Identification,

ITILE OF INVENTION: Nucleic Acid Molecules and Therapy of Ovarian Cancer

ITILE OF INVENTION: Mucleic Acid Molecules and Therapy of Ovarian Cancer

PILE REFERENCE: MRI-030

CURRENT APPLICATION NUMBER: US/10/97,340

CURRENT APPLICATION NUMBER: 60/276,025

PRIOR APPLICATION NUMBER: 60/276,025

PRIOR APPLICATION NUMBER: 60/35,149

PRIOR APPLICATION NUMBER: 60/35,149

PRIOR APPLICATION NUMBER: 60/35,149

PRIOR APPLICATION NUMBER: 60/35,149

PRIOR APPLICATION NUMBER: 60/31,732

PRIOR APPLICATION NUMBER: 60/325,102

PRIOR APPLICATION NUMBER: 60/323,580

PRIOR APPLIC
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; Sequence 1, Application US/10097340

; Publication No. US20030087250A1
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-072-621-2
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                                                                                                                                        Best Loc
Matches
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NUMBER OF SEQ ID NOS: 10
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2
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APPLICANT: Manjula GANN
APPLICANT: Sebastian HC
                                                                                                                                     / Match 100.0%; Score 25; DB 1 Local Similarity 100.0%; Pred. No. 0.07; Pred. No. 0.07; Mismatches
1065 AGCTTCCAACCACGTGTAAATCCTA 1041
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Shubhangi KAMATKAR
Steve G. KOVATS
Rachel E. MEYERS
Michael MORRISEY
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Robert C. BAST, Jr.
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26-OCT-2000; 2000US-0243542P

(BGHM) BRIGHAM & WOMENS HOSPITAL INC.

25-OCT-2001; 2001WO-US049856

02-MAY-2002.

Treating a subject (at risk of) having a hematologic malignancy or multidrug resistance, e.g. lymphoma or myeloma, by administering hypoxia inducible factor 1 binding molecules or small ubiquitin-like-modifier-1 binding molecules.

WPI; 2002-471427/50.

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ALIGNMENTS

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ID AAD390
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XX Human 1
XX Human;
XW Human;
XW Hypoxi
XW Hypoxi
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XX Key
FT Misc_b
FT Misc_b
FT Wisc_b
FT Wisc
PN WO2002
XX Treatin
PT Treatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; haematologic malignancy; multidrug resistance; MDR; SUMO-1; hyppoxia inducible factor-1; small ubiquitin-like-modifier; HIF-1; lymphotold disorder; chronic lymphoproliferative disorder; lymphoma; myeloid disorder; lymphocytic leukaemia; thrombocythaemia; myeloma; angiogenic myeloid metaplasia; myeloid leukaemia; gene therapy; polycythaemia vera; hypoxia responsive element; HRE; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human mdrl gene HIF-1 binding site DNA #4.
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          Treating a subject (at risk of) having a hematologic malignancy or multidrug resistance, e.g. lymphoma or myeloma, by administering hypoxia inducible factor 1 binding molecules or small ubiquitin-like-modifier-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; haematologic malignancy; multidrug resistance; MDR; SUMO-1; hypoxia inducible factor-1; small ubiquitin-11ke-modifier; HIF-1; lymphoid disorder; chronic lymphopoliferative disorder; lymphoma; myeloid disorder; lymphocytic leukaemia; thrombocythaemia; myeloma; angiogenic myeloid metaplasia; myeloid leukaemii; gene therapy; polycythaemia vera; hypoxia responsive element; HRE; antisense;
                                                                             WPI; 2002-471427/50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               or HIF-1-SUMO-1 complex modulators. mdr1-hypoxia responsive
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Pred. No. 0.025;
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RESULT 3
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ABK87742;
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AC ABK87742;
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DE Multiple drug resist XX
DE MURL; ds; multiple d XX
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   The invention relates to a primer for amplifying a nucleic acid comprising a 15 to 30 base continuous oligonuclectides of an MDR1 (multiple drug resistance gene 1) gene fragment appearing as ABK87740. Also included are a kit for quantitatively determining the expression level of MDR1 gene, a probe for capturing a nucleic acid, a reagent fo amplifying a nucleic acid and diagnosis of the drug resistance of a patient. The primer is useful for the determination of MDR expression level by nucleic acid amplification. The present sequence is a fragmen
                                                                                                                                                                                                                                                                                                                                                                                    A primer for amplifying a nucleic acid comprises a specific oligonucleotide sequence for determination of Multiple Drug gene expression level by nucleic acid amplification.
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Pred. No. 0.025;
0; Mismatches (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a primer for amplifying a nucleic acid comprising a 15 to 30 base continuous oligonucleotides of an MDR1 (multiple drug resistance gene 1) gene fragment appearing as ABR8740. Also included are a kit for quantitatively determining the expression level of MDR1 gene, a probe for capturing a nucleic acid, a reagent for amplifying a nucleic acid and diagnosis of the drug resistance of a patient. The primer is useful for the determination of MDR expression level by nucleic acid amplification. The present sequence is a fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A primer for amplifying a nucleic acid comprises a specific oligonucleotide sequence for determination of Multiple Drug gene expression level by nucleic acid amplification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MDR1; ds; multiple drug resistance gene; nucleic acid amplification; MDR1 expression level; multiple drug resistance.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABK87743 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 9; Page 6; 7pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-552738/59.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABK87743;
                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 120 BP; 44 A; 27 C;
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25; Conserv
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                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                         21 G;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                  Score 25; DB 6
Pred. No. 0.03;
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AAI50383;

AAI50383 standard; DNA; 172 BP

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RRESULT 6
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26-MAY-2000; 2000US-0207456P.
30-JUN-2000; 2000US-00608408.
03-AUG-2000; 2000US-00632366.
21-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0236359P.
04-OCT-2000; 2000GB-00024263.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Probe #19069 used to measure gene expression in human placenta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-OCT-2001
                                                                                                                                                                                                                                      Human; bone microarray;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to single exon nucleic acid probes (SENP). The present sequence is one such probe. The probes are useful for producing a microarray for predicting, measuring and displaying gene expression in samples derived from human placenta. The probes are useful for antenatal diagnosis of human genetic disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Penn
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                                                                                                                                                                                                                                                                                                             Human bone marrow expressed single exon probe SEQ ID NO: 18944.
  04-FEB-2000; 2000US-0180312P
                                               30-JAN-2001;
                                                                                               09-AUG-2001
                                                                                                                                                                                              Homo sapiens.
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l Similarity 100.0%;
25; Conservative 0
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Pred. No.
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                                                                                                                                                                                                                                              lymphoma;
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30-JUN-2000; 2000US-00608408.
03-AUG-2000; 2000US-00632366.
21-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0236359P.
04-OCT-2000; 2000GB-00024263.
          Single
                                                                                                                            04-FEB-2000; 2000US-0180312P.
26-MAY-2000; 2000US-0207456P.
30-JUN-2000; 2000US-00608408.
                             WPI; 2001-483446/52
                                                  Penn
                                                                                                 21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             samples, which may enable the improved diagnosis and treatment of such as lymphoma, leukaemia and myeloma. The present sequence is the probes of the invention
                                                                                                                     03-AUG-2000;
                                                                                                                                                                    30-JAN-2001; 2001WO-US000667
                                                                                                                                                                                        09-AUG-2001.
                                                                                                                                                                                                         WO200157275-A2
                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                       Human; brain expressed exon; gene expression analysis; probe; microarray; Alzheimer's disease; multiple sclerosis; schizophrenia; epilepsy; cancer;
                                                                                                                                                                                                                                                                                       Human
                                                                                                                                                                                                                                                                                                         05-NOV-2001
                                                                                                                                                                                                                                                                                                                            AAK18477;
                                                                                                                                                                                                                                                                                                                                               AAK18477 standard; DNA; 172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 172 BP; 58 A; 43 C; 29 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human
                                                                     (MOLE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           bone marrow.
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                                                                                                                                                                                                                                                                                      brain
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                                                                                                                                                                                                                                                                                                                                                                                               62
          exon nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                 ب
                                                                                                                                                                                                                                                                                                                                                                                                                                     25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              which are derived from genomic sequences expressed in the human arrow. They can be used to measure gene expression in bone marrow s, which may enable the improved diagnosis and treatment of cancers
                                                                     MOLECULAR DYNAMICS
                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                                                                                              AGCTTCCAACCACGTGTAAATCCTA 86
                                                                                                                                                                                                                                                                                                                                                                                                           AGCTTCCAACCACGTGTAAATCCTA 25
                                                  Hanzel
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID
                                                                                     ; 2000US-00632366.
; 2000US-0234687P.
; 2000US-0236359P.
; 2000GB-00024263.
                                                                                                                                                                                                                                                                                    expressed single exon probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            invention provides a number of single exon
                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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            acid
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                                                 Chen W,
          probes
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Pred. No.
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         for analyzing
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          gene expression
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RESULT 8
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Best Local S
Matches 25
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26-MAY-2000;
30-JUN-2000;
The invention relates to a single exon nucleic acid probe (SENP) (I) for measuring human gene expression in a sample derived from human adult liver. comprising one of 13109 defined nucleotide sequences given in the specification (or complements) fragments). The probe hybridises at high stringency to a nucleic acid molecule expressed in the human adult liver. (I) may be used for predicting, measuring and displaying gene expression in samples derived from human adult liver. The genes identified may be involved in genetic liver diseases such as cirrhosis, hyperlipoproteinaemia, hyperlipidaemia and hypercholesterolaemia which is
                                                                                                                                                                                  gene
                                                                                                                                                                                                                                                          Penn
                                                                                                                                                      Claim
                                                                                                                                                                                                                              WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      probes which are derived from genomic sequences expressed in the human brain. They can be used to measure gene expression in brain cell samples, which may enable the diagnosis and improved treatment of nervous system diseases such as Alzheimer's disease, multiple sclerosis, schizophrenia, epilepsy and cancers. The present sequence is one of the probes of the
                                                                                                                                                                                                                                                                                                                  04-OCT-2000;
                                                                                                                                                                                                                                                                                                                                             03-AUG-2000;
21-SEP-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hyperlipoproteinaemia;
coronary heart disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human liver single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200157273-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABS44044;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABS44044 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 172 BP; 58
                                                                                                                                                                                                                                                                                     (MOLE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 4; SEQ ID NO 18468; 650pp + Sequence Listing; English
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                                                                                                                                                                                expression
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                                                                                                                                                                                                                                                        SG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sapiens.
                                                                                                                                                      4.
                                                                                                                                                                             genome-derived single exon nucl
expression in human adult liver.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   single exon nucleic acid probe; liver; cirrhosis;
ipoproteinaemia; hyperlipidaemia; hypercholesterol
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25;
                                                                                                                                                                                                                                                                                     MOLECULAR DYNAMICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                      SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AGCTTCCAACCACGTGTAAATCCTA 25
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                                                                                                                                                                                                                                                        Hanzel DK,
                                                                                                                                                                                                                                                                                                               ; 2000US-0180312P.
2000US-0207456P.
2000US-00608408.
; 2000US-00632366.
; 2000US-0234687P.
; 2000US-0234359P.
; 2000US-0234263.
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ilarity 100.0%;
Conservative (
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                                                                                                                                                   NO 19034; 658pp;
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                                                                                                                                                                                                                                                        Chen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      172
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Pred. No. 0.0
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ
                                                                                                                                                                                                                                                          Rank
                                                                                                                                                   English.
                                                                                                                                                                                                 nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hypercholesterolaemia;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 4;
                                                                                                                                                                                               probes useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     <u>,</u>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Other;
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RESULT 9
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Best Local S
Matches 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hermansky-Pudlak syndrome; sarcoidosis; pulmonary haemosiderosis; pulmonary histiccytosis; lymphangioleiomycmtosis; Karagener syndrome; pulmonary alveolar proteinosis; fibrocystic pulmonary dysplasia; primary ciliary dyskinesis; pulmonary hypertension; hyaline membrane disease; open reading frame; ORF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; ds; single exon probe; asthma; lung cancer; COPD; ILD chronic obstructive pulmonary disease; interstitial lung diseamilial idiopathic pulmonary fibrosis; neurofibromatosis; tuberous sclerosis; Gaucher's disease; Niemann-Pick disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         associated with coronary heart disease. ABS25011-ABS51005 represent human liver single exon nucleic acid probes of the invention. Note: The sequence information for this patent does not appear in the printed specification but was obtained in electronic format directly from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ftp.wipo.int/pub/published_pct_sequences
The invention relates to a spatially-addressable set of single exon nucleic acid probes for measuring gene expression in a sample derived from human lung comprising single exon nucleic acid probes having one of 12614 nucleic acid sequences mentioned in the specification, or their complements or the 12387 open reading frames derived from the 12614 probes. Also included are a microarray comprising the novel set of probes it novel set of probes which hybridise at high stringency to a nucleic acid expressed in the human lung; measuring gene expression in a sample derived from human lung, comprising (a) contacting the array with a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-FEB-2000;
26-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-JAN-2001; 2001WO-US000665
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                                                                                                                                                                                                                                                                                                     Claim 4;
                                                                                                                                                                                                                                                                                                                                                         Spatially-addressable set of single exon nucleic measure gene expression in human lung samples.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-JUN<sup>2</sup>2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200186003-A2
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                                                                                                                                                                                                                                                                                                     SEQ ID NO 18613; 634pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; 2000US-0180312P.
; 2000US-0207456P.
; 2000US-00608408.
; 2000US-00632366.
; 2000US-0234687P.
; 2000US-0236359P.
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100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chen W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rank DR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               42 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.032;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 172
                                                                                                                                                                                                                                                                                                                                                                                           acid probes,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ID No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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cc algorithmically predicting at least one exon from problem. Cc the eukaryote; and (b) detectably cc labeled nucleic acids from eukaryote lung mRNA, to a single exon probe, cc in the above mentioned microarray; assigning exons to a single gene, cc in the above mentioned microarray; assigning exons to a single gene, cc comprising (a) identifying exons from genomic sequence by the method cc comprising (a) identifying exons from genomic sequence by the method cc comprising (a) identifying exons from genomic sequence by the method cc above and (b) measuring the expression of each of the exons in several ctissues and/or cell types using hybridisation to a single gene, cc expression of the exons in the tissues and/or cell types indicates that cc the exons should be assigned to a single gene; a peptide comprising one cc filll sequences, mentioned in the specification, or encoded by the exons should be assigned to a single gene; a peptide comprising one cc probes/open reading frames (ORF). The probes are used for gene expression cc analysis, and for identifying exons in a gene, particularly using human cc disease (ILD), familial idiopathic pulmonary disease such as asthma, lung cdisease (ILD), familial idiopathic pulmonary fibrosis, neurofibromatosis, tuberous sclerosis, Gaucher's disease, Niemann-pick disease, Hermansky-pulmonary disease (COPD), interstitial lung cc disease (ILD), familial idiopathic pulmonary haemosiderosis, pulmonary cc interstitial proteinosis, springener syndrome, fibrocystic pulmonary haemosiderosis, pulmonary cc dyskinesis, pulmonary haemosiderosis, pulmonary cc interstitial proteinosis, the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              collection of detectably labeled nucleic acids derived from human mRNA, and (b) measuring the label detectably bound to each probe carray; identifying exons in a eukaryotic genome, comprising (a) algorithmically predicting at least one exon from genomic sequence
             Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequences of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      m human lung
probe of the
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Sequence 172 BP; 58 A; 43 C; 29 G; 42 T; 0 U; 0

Matches Query Match Best Local : 25; Similarity Conservative 100.0%; 100.0%; 0; Pred. Score 25; Mismatches No. 0.032; DB 6; Length 172; Indels 0 0

밁 S 82 Н AGCTTCCAACCACGTGTAAATCCTA 25 AGCTTCCAACCACGTGTAAATCCTA 86

RESULT 10 ABZ35172/c ABZ35172 standard; cDNA;

05-FEB-2003 (first entry)

Human gene expression profile polynucleotide SEQ Ħ ö

bronchial epithelium; prostate; muscle; lung fil tumour; microarray; genome mapping; antibiotic; expression; artery; endothelium; umbilical; vein; aorta; pulmonary artery;
ial epithelium; prostate; muscle; lung fibroblast; osteoblast;
; microarray; genome mapping; antibiotic; antiviral; antifungal gene; antifungal;

Homo sapiens.

WO200274979-A2

20-MAR-2002; 2002WO-US008456

20-MAR-2001; 2001US-0276947P

(ORTH) ORTHO CLINICAL DIAGNOSTICS INC

WPI; 2002-740862/80

New gene expression profile generated from primary, endothelial.

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RESULT 11
AAK45086/c
ID AAK450
XX AAK450
XX AAK450
XX Human;
KW Human;
KW microa
XX Homo s
PN WO2001
XX WO2001
XX Homo s
PN WO2001
XX Homo s
PN WO2001
XX Homo s
XX Homo s
PN WO2001
XX Homo s
XX Homo s
PN WO2001
XX Homo s
PN WO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 general types (NAZZJABNY-ANZZJABNY) and generated from a cell type. The cell type of is a coronary artery endothelium, umbilical artery or vein endothelium, coronated the covascular endothelium, pulmonary artery cendothelium, myometrium microvascular endothelium, pulmonary artery cendothelium, myometrium microvascular endothelium, keratinocyte epithelium, renal cortical epithelium, renal proximal tubule epithelium, cenal airway epithelium, renal epithelium, renal proximal tubule epithelium, cenal airway epithelium, renal epithelium, mammary epithelium, prostate epithelium, cenal proximal tubule epithelium, cenal proximal muscle, astrocytes, cenal fibroblast, cenal proximal cells, coronary artery smooth muscle, central amooth muscle, uterine smooth muscle, lung fibroblast, central proximal cells, ceronary artery smooth muscle, central tubule epithelium, cells, skeletal muscle, seed for determining the level of RNA expression for a sample, determining the proximal cell types. The gene or a protein expression profiles may provide meaningful information with respect to tumour type and stage, central proximal cells, central proximal cells, central proximal cells, central protein expression profile may also be used for creating microarrays. The microarray is useful for genetic and physical mapping of genomes, DNA sequencing, genetic or dedical diagnosis, genotyping of organisms, confirming cell or tissue identifications and in identifying promising antibiotics, antiviral or cells, served to tubour type and stage.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            밁
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Best Local S
Matches 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; bone marrow expressed exon; microarray; cancer; leukaemia; lym;
                                                                                                                                                                                                                                                                                                 04-FEB-2000;
26-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                            30-JAN-2001;
                                                                                                                                                                                                                                      03-AUG-2000;
                                                                                                                                                                                                                                                                   30-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200157276-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAK45086
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAK45086 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention relates to a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            204
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                                                                                 MOLECULAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGCTTCCAACCACGTGTAAATCCTA 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            209
                Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             marrow
                                                                                                                                                               2000US-0180312P.
2000US-0207456P.
2000US-00608408.
2000US-00632366.
2000US-0234687P.
2000US-0234687P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP;
                                                                                                                                                                                                                                                                                                                                                                                               2001WO-US000668
                                                                                                                                           2000GB-00024263.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         65 A; 39
                                                                             DYNAMICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         expressed exon; gene expression analysis; probe; leukaemia; lymphoma; myeloma; ss.
                   Chen W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ç
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             a gene expression profile comprising one or more and generated from a cell type. The cell type
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    single exon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 25;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            67
                   뮸
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.032;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0 U; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 209;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19643.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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RESULT 12
AAK19129/c
ID AAK191
XX AAK191
XX AAK191
XX Human,
XX Human,
XX Human,
XX Human,
XX Human,
XX Homo 8
XX WO2001
XX WO2001
XX O4-FEE
PR 26-MAY
PR 30-JUN
PR 03-JUN
PR 03-JUN
PR 03-JUN
PR 27-SEE
PR 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         먉
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                   04-FEB-2000; 2000US-0180312P.
26-MAY-2000; 2000US-0207456P.
30-JUN-2000; 2000US-00608408.
03-AUG-2000; 2000US-00632366.
21-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-02346359.
04-OCT-2000; 2000GB-00024263.
                                                                                                                                                                                                                                  Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          probes which are derived from genomic sequences expressed in the human bone marrow. They can be used to measure gene expression in bone marrow samples, which may enable the improved diagnosis and treatment of cances such as lymphoma, leukaemia and myeloma. The present sequence is one of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-JAN-2001; 2001WO-US000667.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAK19129;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAK19129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 219 BP; 54 A; 38 C;
                                                                                                                                                                                                                                                                                                                                                                          (MOLE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200157275-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human brain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the probes of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example
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                                                                                                                                                                                                                                  exon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ب
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        brain
                                                                                                                                                                                                                                                                                                                                                                          MOLECULAR DYNAMICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGCTTCCAACCACGTGTAAATCCTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AGCTTCCAACCACGTGTAAATCCTA 119
                                                                                                                                                                                                                                  nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                a expressed exon; gene expression analysis; probe; microarray;
disease; multiple sclerosis; schizophrenia; epilepsy; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 expressed single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ID NO 19643; 658pp + Sequence Listing;
                                                                                                                                                                                                                                                                                                                             Ľ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
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                                                                                                                                                                                                                                                                                                                             Chen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    provides a number of single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           219 BP
                                                                                                                                                                                                                                  probes
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                                                                                                                                                                                                                                                                                                                             Σ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       51 G; 76 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 25; I
Pred. No. 0.
                                                                                                                                                                                                                                for analyzing
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                                                                                                                                                                                                                                                                                                                             뮸
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .033;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ID NO: 19120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  English
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The present invention provides a number of single exon nucleic acid probes which are derived from genomic sequences expressed in the human brain. They can be used to measure gene expression in brain cell samples, which may enable the diagnosis and improved treatment of nervous system diseases such as Alzheimer's disease, multiple sclerosis, schizophrenia, epilepsy and cancers. The present sequence is one of the probes of the

Example 4;

SEQ

ID NO

19120;

650pp +

Sequence

Listing;

English

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ABS19335/c
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ID ABS19335/c
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-FEB-2000;
26-MAY-2000;
30-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hermansky-Pudlak syndrome; sarcoidosis; pulmonary haemosiderosis; pulmonary histiccytosis; lymphangioleiomycmtosis; Karagener syndrome; pulmonary alveolar proteinosis; fibrocystic pulmonary dysplasia; primary ciliary dyskinesis; pulmonary hypertension; hyaline membrane disease; open reading frame; ORF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; ds; single exon probe; asthma; lung cancer; COPD; ILD chronic obstructive pulmonary disease; interstitial lung disease; familial idiopathic pulmonary fibrosis; neurofibromatosis; tuberous sclerosis; Gaucher's disease; Niemann-Pick disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                invention
                           The invention relates to a spatially-addressable set of single exon nucleic acid probes for measuring gene expression in a sample derived from human lung comprising single exon nucleic acid probes having one of 12614 nucleic acid sequences mentioned in the specification, or their complements or the 12387 open reading frames derived from the 12614 probes. Also included are a microarray comprising the novel set of probes in the novel set of probes which hybridise at high stringency to a nucleic acid expressed in the human lung; measuring gene expression in a sample derived from human lung; comprising (a) contacting the array with a collection of detectably labeled nucleic acids derived from human lung (a) contacting the probe of the measuring the label detectably bound to each probe of the array, identifying exons in a sukaryotic genome, comprising (a)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-JAN-2001; 2001WO-US000665
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-NOV-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                Claim 4; SEQ ID NO 19326; 634pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Spatially-addressable set of single exon nucleic acid probes, measure gene expression in human lung samples.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-114183/15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
      algorithmically predicting at least one exon
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       genome-derived single exon probe ORF from lung SEQ ID No 19326.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 AGCTTCCAACCACGTGTAAATCCTA 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULAR DYNAMICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AGCTTCCAACCACGTGTAAATCCTA 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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2000US-00632366.
2000US-0234687P.
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2000US-0207456P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 25; 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chen W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ü
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rank
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         76 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.033;
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      from genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ILD;
      sequences
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RESULT 14
AAI37193
ID AAI37193
AC AAI37
XX AAI37
AC AAI37
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Best Local
                                                                                                                                                                                                                                                                                                                                                                         04-FEB-2000;
26-MAY-2000;
30-JUN-2000;
                                                                                                                                                                                                                                                                                    03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 219 BP; 54 A; 38 C; 51 G; 76 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Probe; microarray; human; placenta; antenatal diagnosis; genetic disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAI37193 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-JAN-2001; 2001WO-US000663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200157272-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Probe #5879 used to measure gene expression in human placenta sample
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      143
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                                                                                                                                                                                              MOLECULAR DYNAMICS INC
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                                                                                                                             Hanzel
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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2000US-00632366.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  2000US-0180312P
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2000US-0236359P.
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                                                                                                                             DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA; 463
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                                                                                                                                Chen W,
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Pred.
                                                                                                                                Rank DR
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Human genome-derived single exon nucleic acid probes useful for analyzing

WPI; 2001-488897/53.

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ID ARX311296
ARX ARX31124
AC AAX311
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26-MAY-2000; 2000US-0207456P.
30-JUN-2000; 2000US-00608408.
03-AUG-2000; 2000US-00632366.
21-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0236359P.
04-OCT-2000; 2000GB-00024263.
                                                                                                                                                                               The present invention provides a number of single exon nucleic acid probes which are derived from genomic sequences expressed in the human bone marrow. They can be used to measure gene expression in bone marrow samples, which may enable the improved diagnosis and treatment of cancer such as lymphoma, leukaemia and myeloma. The present sequence is one of the invention
                                                                                                                            Sequence 463
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human genome-derived single exon nucleic acid probes useful for analyzing gene expression in human bone marrow.
                                                                                                                                                                                                                                                                                                                                                                                           Example 4; SEQ ID NO 5853; 658pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-488900/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to single exon nucleic acid probes (SENP). The present sequence is one such probe. The probes are useful for producing a microarray for predicting, measuring and displaying gene expression in samples derived from human placenta. The probes are useful for antenatal diagnosis of human genetic disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-JAN-2001; 2001WO-US000668
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ch 100.0%; Score 25; DB 4; Length 463; Similarity 100.0%; Pred. No. 0.036; 25; Conservative 0; Mismatches 0; Indels
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BH696214 BOMC13BTF
BH696214 BOMC18TF
BH698271 BOMC18TF
AZ341385 IMO073B15
CC951154 BOIGM45TF
AZ941385 IMO073B15
CC951154 BOIGM45TF
BH983322 Oct602c02.
BZ010051 Oct74007.
CC967311 BOIHM63TR
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ALIGNMENTS

| Qy | Query Match
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| 1 AGCTTCCAACCACGTGTAAATCCTA 25 | Query Match 100.0%; Score 25; DB 9; Length 3843;
Best Local Similarity 100.0%; Pred. No. 0.28;
Matches 25; Conservative 0; Mismatches 0; Indels 0; Gaps 0; | /mol_type="genomic DNA" /db_xref="taxon:9606" <1>3843 /gene="ABCB1" /locus_tag="HCM3396" | | Clark, A.G., Glanowski, S., Nielson, R., Thomas, P., Kejariwal, A., Todd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F., Murphy, B., Todd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F., Murphy, B., Perriera, S., Wang, G., Zheng, X.H., White, T.J., Sninsky, J.J., Adams, M.D. and Cargill, M. Direct Submission Submitted (16-NOV-2003) Celera Genomics, 45 West Gude Drive, Rockville, MD 20850, USA | 1 (bases 1 to 3843) Clark,A.G., Glanowski,S., Nielson,R., Thomas,P., Kejariwal,A., Clark,A.G., Glanowski,S., Nielson,R., Lu,F., Murphy,B., Todd,M.A., Tanenbaum,D.M., Civello,D.R., Lu,F., Murphy,B., Ferriera,S., Wang,G., Zheng,X.H., White,T.J., Sninsky,J.J., Adams,M.D. and Cargill,M. Inferring nonneutral evolution from human-chimp-mouse orthologous gene trios gene trios Science 302 (5652), 1960-1963 (2003) | AY408954 3843 bp DNA linear GSS 15-DEC-2003 Homo sapiens ABCB1 gene, VIRTUAL TRANSCRIPT, partial sequence, AY408954 AY408954 AY408954.1 GI:39764922 GSS. Homo sapiens (human) Homo sapiens (human) Enkaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. |

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3182 bp mRNA linear HTC 03-APR-20 MNB musculus 11 days pregnant adult female ovary and uterus cDNA, RIKEN full-length enriched library, clone:5031438E12 product:ATP-binding cassette, sub-family B (MDR/TAP), member 1B, full insert sequence.
                                 Carninci, P. and Hayashizaki, Y.
High-efficiency full-length cDNA cloning
Meth. Enzymol. 303, 19-44 (1999)
                                                                                                   Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Clark, A.G., Glanowski, S., Nielson, R., Thomas, P., Kejariwal, Todd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F., Murphy, B., Ferriera, S., Wang, G., Zheng, X.H., White, T.J., Sninsky, J.J., Adams, M.D. and Cargill, M.
                                                                                                                                                                                                 AK030328.1 GI:26081769
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (16-NOV-2003) Celera Genomics, Rockville, MD 20850, USA
This sequence was made by sequencing geno
                                                                                                                                                             Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Clark, A.G., Glanowski, S., Nielson, R., Thomas, P., Kejariwal, A., Todd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F., Murphy, B., Ferriera, S., Wang, G., Zheng, X.H., White, T.J., Sninsky, J.J., Adams, M.D. and Cargill, M.
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Mammalia; Eutheria; Primates;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /gene="ABCB1"
/locus_tag="HCM3396"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 organism="Pan troglodytes"
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100.0%; Pred. No. 0.
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URL:http://fantom.gsc.riken.jp/.
Location/Qualifiers
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                                                                                    /db_xref="FANTOM_DB:5031438E12"
/db_xref="taxon:10090"
                                                                                                                                                        organism="Mus musculus"
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ALL Nature 420, 563-573 (2002)

86 (bases 1 to 3182)

88 Adachi, J., Aizawa, K., Akimura, T., Hara, A., Hashizume, W., Fukuda, S., Furumo, M., Hanagaki, T., Hara, A., Hashizume, W., Fukuda, S., Furumo, M., Hiramoto, K., Hiraoka, T., Hirozane, T., Hayashida, K., Hayatsu, N., Hiramoto, K., Kagawa, I., Kasukawa, T., Kotoh, H., Kawai, J., Kojima, Y., Kondo, S., Konno, H., Kouda, M., Koya, S., Kurihara, C., Matsuyama, T., Miyazaki, A., Murata, M., Nishi, K., Nomura, K., Numazaki, R., Ohno, M., Ohsato, N., Nakamura, M., Nishi, K., Nomura, K., Numazaki, R., Ohno, M., Ohsato, N., Saito, R., Saito, H., Sakai, C., Sakai, K., Sakazume, N., Okazaki, Y., Sasto, R., Saito, H., Sakai, C., Sakai, K., Sakazume, N., Sagabe, Y., Tagami, M., Tagawa, A., Takahashi, F., Takaku-Akahira, S., Takada, Y., Tanaka, T., Toman, A., Toya, T., Yasunishi, A., Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  submitted (16-JUL-2001) Yoshihide Hayashizaki, The Institute of Physical and Chemical Research (RIKEN), Laboratory for Genome Exploration Research Group, RIKEN Genomic Sciences Center (GSC), RIKEN Yokohama Institute; 1-7-22 Suehiro-cho, Tsurumi-ku, Yokoha Kanagawa 230-0045, Japan (E-mail:genome-res@gsc.riken.jp, URL:http://genome.gsc.riken.jp/, Tel:81-45-503-9222, Pax:81-45-503-9216)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CDNA library was prepared and sequenced in Mouse Genome
Encyclopedia Project of Genome Exploration Research Group
Genomic Sciences Center and Genome Science Laboratory in R
Division of Experimental Animal Research in Riken contribu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Group Phase I & II Team.
Analysis of the mouse transcriptome based on functional annotation of 60,770 full-length cDNAs
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/tissue_type="ovary and uterus"
/clone_Tib="RIKEN full-length enriched mouse
/dev_stage="11 days pregnant adult"
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Euteleostomi; Sus.

Vector identified

with

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Best Local Similarity
Matches 23; Conserv
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CF176979.1 GI:33288755
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Sus scrofa (pig)
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Clark, A.G., Glanowski, S., Nielson, R., Thomas, P., Kejariwal, A., Todd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F., Murphy, B., Ferriera, S., Wang, G., Zheng, X.H., White, T.J., Sninsky, J.J., Adams, M.D. and Cargill, M.
Inferring nonneutral evolution from human-chimp-mouse orthologous
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                                                                                                                                                                                                        Submitted (16-NOV-2003) Celera Genomics, Rockville, MD 20850, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Clark, A.G., Glanowski, S., Nielson, R., Thomas, P., Kejariwal, A., Todd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F., Murphy, B., Todd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F., Murphy, B., Ferriera, S., Wang, G., Zheng, X.H., White, T.J., Sninsky, J.J., Adams, M.D. and Cargill, M.

Direct Submission
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                        AGCTTCCAGCCACGGGTAAATCCTA 605
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based on alignment.
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                                                                                                                                                                                                                                                                                                                                                          /organism="Mus musculus"
/mol_type="genomic DNA"
/db_xref="taxon:10090"
<1...>3816
                                                                                                                                                                                                                                                                                                                    'gene="ABCB1"
'locus_tag="HCM3396"
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Best Local Similarity 91.7%;
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Smith, T. P.L., Freking, B.A., Ford, J.J., Vallet, J.L., Fox, J., wise, T.A., Nonneman, D.J., Wray, J.E. and Keele, J.W. A second set of porcine ESTs from a pooled-tissue normalized
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: smith@email.marc.usda.gov
Single pass sequencing. Bases called with phr
trimmed with the aid of the trim_alt option.
cross match v0.990329.
plate: SRG8002 row: F column: 24
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Contact: Smith TPL
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                                                                                                      FBN Dummerstorf
2nd wilhelm-Stahl-Allee, Dummerstor
Tel: 0049-038208-68700
Fax: 0049-038208-68702
Email: schwerin@fbn-dummerstorf.de
                                                                                                                                                                                                                                                                                                                                                              EST.
Bos taurus (cow)
Bos taurus
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fbn-i094 Cattle pUC18 Library
BE217416
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Fax: 402 762 4390
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PO Box 166, Clay Center, N
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Schwerin, M., Dorroc
                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                           Unpublished (2000)
Contact: Schwerin M
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                                                                                                                                                                                        Molecular Biology
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                                                                        POLYA=Yes.
                                                                                      High quality sequence
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/lab host="DH10B"
/lab host="DH10B"
/clone lib="MARC 3PIG"
/note="Vector: pcDNA3.1; Site 1: EcoRI; Site 2: NotI;
/notes made with RNA pooled from multiple tissues
including brain, liver, muscle, placenta/endometrium,
ovary, testes, and bone marrow."
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/mol_type="mRNA"

/db_xref="taxon:9823"
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 /organism="Bos taurus"
/mol_type="mRNA"
                                                      Location/Qualifiers
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Pred. No. 28;
0; Mismatches
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CNSOA1N4/c
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Matches
Query Match
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                            Submitted (18-NOV-2003) Genoscope - Centre National de Sequencage :
BP 191 91006 EVRY cedex - FRANCE (E-mail : seqref@genoscope.cns.fr)

The sequences are based on single pass reads.

Life Technologies (a division of Invitrogen) members carried out full-length librairies construction : Temple G.

Genoscope members carried out sequencing and annotation : Castelli V., Aury J.M., Jaillon O., Wincker P., Menard M., Cruaud C., Schachter V., Weissenbach J., Salanoubat M.

URGV INRA : Clepet C., Caboche M.

Annotation is based on the June 2003 version of the Arabidopsis genome released by MIPS (Munich Information center for Protein Sequences). 5 prime and 3 prime are assembled with Phrap.

http://www.genoscope.cns.fr/externe/sequences/Banque_Projet_EF/Full
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Castelli, V., Aury, J.M., Jaillon, O., Wincker, P., Clepet, C.,
Menard, M., Cruaud, C., Quetier, F., Scarpelli, C., Schachter, V.,
Temple, G., Caboche, M., Weissenbach, J. and Salanoubat, M.
Whole Genome Sequence Comparisons and 'Full-Length' cDNA Sequences:
A Combined Approach to Evaluate and Improve Arabidopsis Genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosida; eurosida; Brassicales; Brassicaceae; Arabidopsis.
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806 bp mRNA linear HTC 06-FEB-200
Arabidopsis thaliana Full-length cDNA-Complete sequence from clone
GSLTSIL25ZG04 of Silique of strain col-0 of Arabidopsis thaliana
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Location/Qualifiers
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Arabidopsis thaliana
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                                                                      /gene="At5g51440"
                                                                                                                 /plasmid="pCMVSPORT
                                                                                                                                             /clone="GSLTSIL25ZG04"
/tissue_type="Silique"
                                                                                                                                                                                                                       mol_type="mRNA"
strain="Col-0"
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/note="Vector: pUC18 Sma I/BAP"
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/dev_stage="lactation"
/lab_host="XL1-blue"
                                                                                                                                                                                             db_xref="taxon:3702"
                                                                                                                                                                                                                                                                  organism="Arabidopsis thaliana"
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Score 20.2;
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Seg primer: CACACAGGAAACAGCTATGACC
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489 bp DNA linear GSS 2700224F07R Mouse 10kb plasmid UUGC2M library Mus musculus clone UUGC2M0224F07 R, genomic survey sequence.
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Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             plasmid inserts
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Insert Length: 10000 Std Erro
Plate: 0224 row: F column: 0
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               polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel elettrophoresis. Vector DNA was prepared from a derivative of phota? (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XI10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                            was blunt end-repaired with T4 DNA polymerase and T4
                                                                                                                                                                                                                                                                                                                                                  (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA
                                                                                                                                                                                                                                                                                                                                                                                                                              musculus C57BL/6J (female) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /lab_host="E. coli strain XL10-Gold, T1-resistant, F
/clone_lib="Mouse 10kb plasmid UUGC2M library"
/note="Vector: FWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sex="Female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  organism="Mus musculus"
/mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ocation/Qualifiers
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603307314F1 NCI_CGAP_Mam6 Mus
mRNA sequence.
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cDNA Library Preparation: Life Technogies, Inc.

cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)

DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information can
                                                                          BH521842
BH521842.1 GI:17729927
GSS.
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National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BI69163
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mus musculus (house mouse)
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Brassica oleracea
Brassica oleracea
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
                                                                                                                                                                    BH521842 526 bp DNA linear GSS BOGQF60TF BOGQ Brassica oleracea genomic clone BOGQF60,
                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone="IMAGE:534383"
/clone="IMAGE:534383"
/tissue_type="infiltrating ductal carcinoma"
/tissue_type="infiltrating ductal carcinoma"
/tissue_type="infiltrating ductal carcinoma"
/tissue_type="infiltrating ductal carcinoma"
/dev stage="5 months"
/lab_host="DH10B"
/clone=lib="NCI_CGAP_Mam6"

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strain="FVB/N"
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Pred. No. 90;
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: cdtown@tigr.org
DNA is from a doubled haploid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tel: 301-838-3523
Fax: 301-838-0208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       9712 Medical Center Drive,
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                                                                                                                                                                                                   Fax: 81-45-503-9573
                                                                                                                                                                                                                           Tel: 81-45-503-9605
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/strain="TO1000DH3"
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              /organism="Zinnia elegans"
/mol_type="mRNA"
/cultivar="Canary bird"
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                                                                                                     Location/Qualifiers
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Pred. No. 1.7e+02;
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1 (bases 1 to 526)
1 (bases 1 to 526)
Town,C.D., Van Aken,S., Utterback,T., Koo,H. and Fraser,C.M.
                                                                                                                                                             visualization by comprehensive microarray analysis of gene expression programs during transdifferentiation of mesophyll cells into xylem cells eroc. Natl. Acad. Sci. U.S.A. 99 (24), 15794-15799 (2002) Contact: Taku Demura Morphogenesis Research Group RIKEN Plant Science Center 1-7-22 Suehirocho, Yokohama, Kanagawa 230-0045, Japan 1-7-22 Suehirocho, Yokohama, Kanagawa 230-0045, Japan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        553 bp mRNA linear AU291400 zinnia cultured mesophyll cell equalized elegans cDNA clone Z6035, mRNA sequence.
AU291400
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Demura, T., Tashiro, G., Horiguchi, G., Kishimoto, N., Kubo, M., Matsuoka, N., Minami, A., Nagata-Hiwatashi, M., Nakamura, K., Okamura, Y., Sassa, N., Suzuki, S., Yazaki, J., Kikuchi, S. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Zinnia elegans
Zinnia elegans
Zinnia elegans
Zinnia elegans
Zinnia elegans
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
asterids; campanulids; Asterales; Asteraceae; Asteroideae;
Heliantheae; Zinnia.
1 (bases 1 to 553)
1 (bases 1 to 553)
Email: demura@postman.riken.go.jp
This clone was obtained at our laboratory.
Seq primer: M13 forward.
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/note="Vector: pHOS1; Site_1: BstXI; 2-3 kb sheared genomic DNA inserted into pHOS1 using BstXI linkers"
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RESULT 13
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Schalkwyk,L.C., Cusack,B., Dunkel,I., Hopp,M., Kramer,M., Palczewski,S., Piefke,J., Scheel,S., Weiher,M., Wenske,G., Lehrach,H. and Himmelbauer,H.
Advanced integrated mouse YAC map including BAC framework Genome Res. 11 (12), 2142-2150 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Department of Vertebrate Genomics/ Abtlg. Lehrach
Max-Planck-Institute of molecular Genetics
Ihnestr.73, D-14195 Berlin-Dahlem, Germany
Email: himmelbauer@molgen.mpg.de
Information on marker localization (Chromosome, map segment) and
YACs positive for this probe can be obtained from our web site
http://www.molgen.mpg.de/(tilde)rodent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bir51B20 Mouse IRS-PCR fragment
genomic clone Bir51B20, genomic
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Mammalia; Eutheria; Rodentia;
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                                                                                                                                                                                                                                                                        /clone="Bir51B20"
/lab_host="E.coli DH5alpha"
/clone lib="Mouse IRS-PCR fragment library 57R/bir"
/clone lib="Mouse IRS-PCR fragment library was
/note="Vector: pAMP10 (Gibco-BRL); The library was
generated from IRS-PCR fragments obtained by amplification
of genomic mouse DNA with the mouse B1-repeat primer B1R
(5'-AGTTCCAGACAGGGGCTAYACAGA-3'). Additional
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/clone Tib="zinnia cultured mesophyll cell equalized cDNA"
/note="Vector: pGEM-T easy; cultured in tracheary element
differentiation-inductive medium"
                                                                                                                                                                                                                                       information online at
www.molgen.mpg.de/~rodent/html/introduction.html"
                                                                                                                                                                                                                                                                                                                                                                                                                                        /mol_type="genomic DNA"
/strain="C57BL/6"
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/clone="Z6035"
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Pred. No. 1.7e+02;
0; Mismatches 3;
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Pred. No. 1.7e+02;
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BH713924
BH713924.1 GI:18806913
GSS.
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Brassica oleracea
Brassica oleracea
Brassica oleracea
Brassica; Streptophyta; Embryophyta; Tracheophyta;
Eukaryota; Vagnoliophyta; eudicotyledons; core eudicots;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
rossids; eurossids II; Brassicales; Brassicaceae; Brassica.
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                                                          9712 Medical Center Drive,
Tel: 301-838-3523
Fax: 301-838-0208
                                                                                                                                                                                                                                                                                                                                                     GSS.
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BOMNB04TR BO_2_3_KB Brassica oleracea genomic clone BOMNB04
                                                                                                                                                        Other_GSSs: BOMNBO4TF
                                                                                                                                                                        1 (bases 1 to 617)
Town,C.D., Van Aken,S., Utterback,T., Koo,H. and Fraser,C.M. Whole genome shotgun sequencing of Brassica oleracea
Unpublished (2001)
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1 (bases 1 to 604)
1 Town,C.D., Van Aken,S., Utterback,T., Koo,H. and Fraser,C.M.
Mhole genome shotgun sequencing of Brassica oleracea
                    Email: cdtown@tigr.org
DNA is from a doubled haploid provided by Tom Osborn.
                                                                                                                                        Contact: Chris Town
                                                                                                                                                                                                                                                                                                                                                                         BH696214.1 GI:18768830
                                                                                                                                                                                                                                                                                                                                                                                                            genomic survey sequence
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Fax: 301-838-0208
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Unpublished (2001)
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seq primer: TR
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DNA is from a doubled haploid provided by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Other_GSSs: BOMCI38TR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AGCTTCCAACCACGTGTAAATCCT 24
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/note="Vector: pHOS1; Site_1: BstXI; 2-3 kb sheared
genomic DNA inserted into pHOS1 using BstXI linkers"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Brassica oleracea"
/mol type="genomic DNA"
/strain="Tol000DH3"
/db_xref="taxon:3712"
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87.5%;
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Pred. No. 1.7e+02;
                                                                                                Rockville,
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Search completed: February 9, 2005, 21:55:40 Job time : 2451.85 secs
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Best Local Similarity 87.5%;
Matches 21; Conservative
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Best Local Similarity 87.5
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1 (bases 1 to 634)

1 (bases 1 to 634)

1 Town,C.D., Van Aken,S., Utterback,T., Koo,H. and Fraser,C.M. Whole genome shotgun sequencing of Brassica oleracea
Unpublished (2001)

Contact: Chris Town
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BOMCI67TF BO_2_3_KB Brassica oleracea genomic clone BOMCI67,
genomic survey sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       9712 Medical Center Drive, Rockville, MD 20850, USA. Tel: 301-838-3523 Fax: 301-838-0208 Email: Cdtown@tigr.org DNA is from a doubled haploid provided by Tom Osborn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Brassica oleracea
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/clone="BOMNB04"
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/clone="Wector: pHOS1; Site 1: BstXI; 2-3 kb sheared
/note="Wector: pHOS1; Site 1: using BstXI linkers"
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/clone=TbendCIG7"
/clone=Tb
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Maximum Match 100%
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Copyright (c) 1993 - 2005 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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| ב | 25 | 100.0 | 25 | σ | AX504310 | AX504310 Sequence |
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გ | 25 | 100.0 | 25 | σ | AX504327 | AX504327 Sequence |
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| | 25 | 100.0 | 3840 | σ | AX481416 | |
| | 25 | 100.0 | 3843 | σ | BD171402 | BD171402 Method to |
| c 10 | 25 | 100.0 | 3860 | σ | AX322787 | AX322787 Sequence |
| | 25 | 100.0 | 3860 | σ | AX322789 | Sequence |
| | 25 | 100.0 | 3988 | σ | BD190394 | BD190394 Phosphati |
| | 25 | 100.0 | 3988 | σ | AR452556 | AR452556 Sequence |
| c 14 | 25 | 100.0 | 3988 | σ | AX024454 | |
| | 25 | 100.0 | 4192 | v | AF016535 | |
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| | 25 | 100.0 | 4264 | σ | AR051650 | AR051650 Sequence |
| | 25 | 100.0 | 4378 | φ | E02326 | E02326 Multidrug r |
| c 19 | 25 | 100.0 | 4553 | σ | CQ716151 | CQ716151 Sequence |
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RESULT 2
AX504327/c
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AX504327 25 bp Sequence 30 from Patent WO0234291. AX504327

DNA

linear

PAT 27-SEP-2002

AX504327.1 GI:23386139

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ORIGIN

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SOURCE ORGANISM

Homo sapiens (human) Homo sapiens

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| 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | • | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 1000 |
| 98472 | 98472 | BIFF | 8630 | 8630 | 8630 | 8630 | 6505 | 4669 | 4669 | 4669 | 4669 | 4669 | 4669 | 4646 | 4646 | 4646 | 4646 | 4646 | 4646 | 4646 | 4646 | 4646 | 4643 | 4643 | |
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| AX707905 | AX /069/5 | AKU286/2 | AXULZIZI | AX012320 | AR306492 | AR306491 | AR028671 | AR405961 | AR363344 | AR203322 | I08557 | AR091275 | AR055785 | HUMMDR1 | AX504298 | AX391099 | AX336708 | AX336420 | AR380622 | 149610 | CQ861565 | BD234195 | AX587788 | AX522070 | |
| AX707905 Sequence | | | | Sequence | Sequenc | | | | | Sequenc | ന | AR091275 Sequence | AR055785 Sequence | M14758 Homo sapien | AX504298 Sequence | | | | | æ | CQ861565 Sequence | | AASS / /88 Sequence | | |

ALIGNMENTS

REFERENCE AUTHORS TITLE RESULT 1 AX504310 LOCUS DEFINITION ACCESSION VERSION VERSION KEYWORDS SOURCE ORGANISM Query Match 100.0%; Score 25; DB 6; Length 25; Best Local Similarity 100.0%; Pred. No. 0.14; Matches 25; Conservative 0; Mismatches 0; Indels JOURNAL source AX504310 25 bp Sequence 13 from Patent W00234291. AX504310 GI:23386128 multiple drug resistance Patent: WO 0234291-A 13 02-MAY-2002; THE BRIGHAM AND WOMEN'S HOSPITAL, INC. Congositions and methods for treating hematologic malignancies and Homo sapiens (human) Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. /organism="Homo sapiens" /mol_type="unassigned DNA" /db_xref="taxon:9606" Location/Qualifiers DNA (sn) linear PAT 27-SEP-2002

PRI 08-JAN-1995

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Heinrich,G. and Kerb,R.
Methods for treatment of cancer using irinotecan
Patent: WO 03013536-A 674 20-FEB-2003;
Epidauros Biotechnologie AG (DE)
Location/Qualifiers
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Seguence 674 from Patent
AX707906
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multiple drug resistance
Patent: WO 0234291-A 30 02-MAY-2002;
THE BRIGHAM AND WOMEN'S HOSPITAL, INC.
Location/Qualifiers
                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Methods for the treatment of cancer with
Patent: WO 03013534-A 674 20-FEB-2003;
Epidauros Biotechnologie AG (DE)
Location/Qualifiers
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Mammalia; Eutheria;
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/db_xref="taxon:9606"
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J. Biol. Chem. 265 (1), 506-514 (1990)
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Chen, C.J., Clark, D.
Roninson, I.B.
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Chin, J.E., Soffir, R., Noonan, K.E., Choi, K. and Roninson, I.B. Structure and expression of the human MDR (P-glycoprotein) 9
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M29432.1 GI:187481
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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        /gene="PGY1"
/note="PGY1,
536. .739
                                                                                    /gene="PGY1"
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/number=12
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/mol_type="unassigned DNA"
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Homo sapiens P-glycoprotein (
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Direct Submission
Submitted (30-NOV-2003) Human Molecular
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Mammalia; Eutheria; Primates;
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                                                                                                                                                                                                                                           /codon_start=1
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DIRTINVRFLREIIGVVSQEPVLFATTIAENIRYGRENVTMDEIEKAVKEANAYDFIM
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/note="PGY1, intron
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/note="P-glycoprotein"
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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/note="transporter"
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join(<16. .126,297. .422,
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Thiruvananthapuram, Kerala
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Trujimoto, H. and Okai, Y.
Trujimoto, H. and Okai, Y.
Direct Submission
Submitted (05-JUN-1999) Hajime Tsujimoto, The University of bepartment of Veterinary Internal Medicine; The University of Tokyo, Bunkyo-ku, Tokyo 113-8657, Japan (E-mail:atsuji@hongo.ecc.u-tokyo.ac.jp, Tel:+81-3-5841-8004, Fax:+81-3-5841-8178)
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Molecular analysis of multidrug resistance :
Am. J. Vet. Res. 61 (9), 1122-1127 (2000)
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Mammalia; Eutheria; Carnivora; Fissipedia; Felidae;
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FKAVHFSYPSRKEVKILKGLMLKVQSGQTVALVGNSGCGKSTTVQLMQRLYDPTOGNV
SIDGQDIRTINVRYLREI IGVVSQEPVLFATTI AENIRYGRENVTMEEI EKAVKEANA
YDFIMKLPNKFDTLVGERGAQLSGGOKORIAIARALVRNPKILLLDEATSALDSESSA
VVQVALDKARKGRTTI VVAHRLSTIRNADVIAGFDDGVIVEKGNHDELMKEEGI VFKL
VTMQTRGNEI EILENAVYESI SEIDALENSPKDSGSSLIRRSTRKSIHASGQDRKLG
VTMQTRGNEI EILENAVYESI SEIDALENSPKDSGSSLIRRSTRKSIHASGQDRKLG
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DUSMFDDFKITTGALTTRLANDAAQVKGAIGSRLAIITONIANLGTGIIISLIYGMQL
TLLLLAIVPIIAIAGALVEAMLSGQALKDKKELEGAKVATEALENFTTVVSLTREQK
FEYNYAQSLQVYPXNSIRKAHIFGISFSITQAMMYFSYNGCFRFGAYLVAHEFMDFQD
VLLVFSAIVFGAMAVGQVSSFAPDYAKAKVSAAHVIMIIEKIDLIDSYSTEGLMPNTL
EGNNTINGKEIKHLANVQMLRAHMGIVGQEFILFDCSIGENIAVGDUSRVVSGEEIVN
AAKEANIHPFIETLPDKYNTRVGDKGTQLSGGQKQRIAIARALVRQPRILLLDEATSA
LDTESEKVVQEALDKAR"
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WFDVHDVGELNTRLTDDVSKINEGIGDKIGMFFQSMATFFIGFIVGFTRGWKLTLVIL
AISPVLGLSAAIWAKILSSFTDKELLAYAKAGAVAEEVLAAIRTVIAFGGQKKELERY
NKNLEEAKRIGIKKAITANISIGIAFLLIYASYALAFWYGTSLVLSHEYSIGQVLTVF
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/organism="Felis catus"
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/--'' feline lymphoma_cell_line"
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/db_xref="GI:6472653"
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BD171402/c
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Mammalia; Sutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 3843)
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Method for predicting
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BD171402
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Mammalia; Eutheria;
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C12N15/09,C12Q1/68,C12N15/00
Method for predicting side effects
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JP 2002223769-A/1
13-AUG-2002
31-JAN-2001 JP 2001024723
                                                                                                                                                                                                                                                                                                                                                                                     for predicting side effects of immunosuppressant
                                                                                                                                                                                                    /organism='Homo sapiens (human)'
                                                                                                                                                        /organism="Homo sapiens"
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/db_xref="taxon:9606"
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Best Local Similarity
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Phosphatidylcholine as a m
BD190394.1 GI:33000133
JP 2002522381-A/1.
Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates; 1 (bases 1 to 3988)
                                                   Homo sapiens (human)
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Method of identifying and/or isolating
Patent: WO 0192877-A 3 06-DEC-2001;
ST. JUDE CHILDREN'S RESEARCH HOSPITAL (
Location/Qualifiers
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Method of identifying and/or isolating
Patent: WO 0192877-A 1 06-DEC-2001;
ST. JUDE CHILDREN'S RESEARCH HOSPITAL (
LOCALION/Qualifiers
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Mammalia; Eutheria; Primates;
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AX322787
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="unassigned DN
/db_xref="taxon:9606"
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/db_xref="taxon:9606"
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                 Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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AR452556
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AX024454
AX024454.1 GI:10184622
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Stremmel,W.
Patent: DE 19857750-A 1 24-FEB-2000;
STREMMEL WOLFGANG (DE)
                                                                                                                                        Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                         Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                  AX024454
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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/mol_type="genomic
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                                                                                           Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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AP016535
AF016535.1 GI:2353263
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2 (bases 1 to 4192)
2 (chen,G., Duran,G.E., Steger,K.A., Lacayo,N.J., Jaffred Chen,G., Duran,G.E., Steger,K.A., Lacayo,N.J., Jaffred Dumontet,C. and Sikic,B.I.
Multidrug-resistant human sarcoma cells with a mutant Multidrug-resistant human sarcoma cells with a mutant P-glycoprotein, altered phenotype, and resistance to J. Biol. Chem. 272 (9), 5974-5982 (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 4192)
Chen,C.J., Chin,J.E., Ueda,K., Clark,D.P., Pastan,I.,
Gottesman,M.M. and Rominson,I.B.
Internal duplication and homology with bacterial transport protein the mdr1 (P-glycoprotein) gene from multidrug-resistant human
                                                                                                                                                                                                                                                                                                                                                                                      Chen.G., Lacayo,N.J., Steger,K.A. Direct Submission Submitted (28-JUL-1997) Medicine, Medicine, Stanford, CA 94306, USA
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/translation="mdlegdrnggakkknffklnnksekdkkekkptvsvfsmfrysn
wldklymvvgtlaaiihgaglplmmlvfgemtdifanagnledlmsnitnrsdindtg
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/db_xref="taxon:9606"
/chromosome="7"
                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                  cell_type="MES-SA/DxP"
                                                                                                                                                                                    gene="mdr1"
                                                                                                                                                                                                                                                   clone="8.1"
                                                                                                                                                                                                                                                                      map="7q21.1"
                                                                                                  note="multidrug-resistance"
                                                                                                                                   gene="mdr1"
                                                                                                                     function="transporter protein"
                                                                                                                                                                    note="multidrug-resistance
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LAISPYLGLSAANWAKILSSFTDKELLAYAKAGAVAEEVLAAIRTYIAFGGÇKELER
YNKNLEEAKRIGIKKAITANISIGAAFELLIYASYALAFWYGTTLVLSGEYSIGQVLTV
FSVLIGAFSVGQASPSIEAFANAGAAYEIFKIINKESJOSYSKSGHKPDNIKGNLE
FRNUFESYPSRKEKKILKGLUKVQSGQTVALVGNSGCGKSTTVQLMQRLYDFTEGNV
SVDGQDIRTINVRFLREIIGVVSQEPVLFATTIAENIRYGRENVTMDEIEKAVKEANA
YDFIMKLPHKPDTLVGERGAQLSGGQKQRIAIARARLYVFRESTA
VVQVALDKARKGATTIVIAHKLSTVRNADVIAGFDDGVIVEKGNALDESSA
VVQVALDKARKGATTIVIAHKLSTVRNADVIAGFDDGVIVEKGNALDELMKEKGIVEKL
VTMQTAGNEVELLENAADESKSEIDALEMSSNDSRSSLIRKRSTRSVRGSQAQDRKLS
TKDEPETRQNSNYLFSLLFLALGJISFITFFLQGFTGKAGEILTKRLRYWFRESMLRQ
DVSWFDDPKNTTGALTTRLANDAAQVKGAIGSRLAVITQNIANLGTGIISFIYGMQL
TLLLLAIVFIIAAJVEIIAGAALJISFITFFLQGFTGKAGEILTKRLRYWFRESMLRQ
DVSWFDDPKNTTGALTTRLANDAAQVKGAIGSRLAVITQNIANLGTGIISFIYGEQK
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| US-09-270-767-11434 | US-09-134-000C-186 | PCT-US94-06284-4 | US-08-227-370-4 | US-09-949-016-13514 | US-09-566-921-7 | US-08-787-091-1 | US-09-949-016-1772 | US-09-016-434-1422 | US-09-949-016-61079 | US-09-762-195-2 | US-09-023-655-1168 | US-09-970-966-175 | US-09-825-294-175 | US-09-713-550-175 | US-09-640-173-175 | US-09-450-105-1 | US-09-120-513-1 |
| | | | | | | | | 1422, | 61079, | | | 1/5 | 175, | 1/5, | 1/5, | 1, AF | ٦. |
| | Sequence | 70.4 945 4 US-09-134-000C-186 Sequence 69.6 616 4 US-09-270-767-11434 Sequence | 71.2 26 5 PCT-US94-06284-4 Sequence 4, 70.4 945 4 US-09-134-000C-186 Sequence 186 69.6 616 4 US-09-270-767-11434 Sequence 11 | 42 17.8 71.2 26 1 US-08-227-370-4 Sequence 43 17.8 71.2 26 5 PCT-US940-66284-4 43 17.6 70.4 945 4 US-09-134-000C-186 Sequence 44 17.6 70.4 945 4 US-09-270-767-11434 Sequence | 41 18.2 72.8 47698 4 US-09-949-016-13514 Sequence 42 17.8 71.2 26 1 US-08-227-370-4 Sequence 43 17.8 71.2 26 5 PCT-US94-06284-4 Sequence 44 17.6 70.4 945 4 US-09-134-000C-186 Sequence 45 17.4 69.6 616 4 US-09-270-767-11434 Sequence | 40 18.2 72.8 8137 4 US-09-566-921-7 Sequence 41 18.2 72.8 47698 4 US-09-949-016-13514 Sequence 42 17.8 71.2 26 1 US-08-227-370-4 Sequence 43 17.8 71.2 26 5 PCT-US94-06284-4 Sequence 44 17.6 70.4 945 4 US-09-134-000C-186 Sequence 45 17.4 69.6 616 4 US-09-270-767-11434 Sequence | 39 18.2 72.8 1446 3 US-08-787-091-1 Sequence 40 18.2 72.8 8137 4 US-09-566-921-7 41 18.2 72.8 47598 4 US-09-949-016-13514 Sequence 41 18.2 72.8 47698 4 US-09-949-016-13514 Sequence 42 17.8 71.2 26 1 US-08-227-370-4 Sequence 43 17.8 71.2 26 5 PCT-US94-05284-4 Sequence 44 17.6 70.4 945 4 US-09-134-000C-186 Sequence 45 17.4 69.6 616 4 US-09-270-767-11434 Sequence | 18.2 72.8 1255 4 US-09-949-016-1772 Sequence 18.2 72.8 1446 3 US-08-787-091-1 Sequence 18.2 72.8 8137 4 US-09-566-921-7 Sequence 18.2 72.8 47698 4 US-09-949-016-13514 Sequence 17.8 71.2 26 1 US-08-227-370-4 Sequence 17.8 71.2 26 5 PCT-US94-06284-4 Sequence 17.6 70.4 945 4 US-09-134-000C-186 Sequence 17.4 69.6 616 4 US-09-270-767-11434 Sequence | 37 18.2 72.8 723 4 US-09-016-434-1422 Sequence 1422, 38 18.2 72.8 1255 4 US-09-949-016-1772 Sequence 1772, 39 18.2 72.8 1446 3 US-08-787-091-1 Sequence 1, App 40 18.2 72.8 8137 4 US-09-566-921-7 Sequence 7, App 41 18.2 72.8 47698 4 US-09-949-016-13514 Sequence 7, App 71.8 71.2 26 1 US-08-27-370-4 Sequence 4, App 71.8 71.2 26 5 PCT-US94-06284-4 Sequence 4, App 71.6 70.4 945 4 US-09-270-767-11434 Sequence 186, 18.5 17.4 69.6 616 4 US-09-270-767-11434 Sequence 11434. | 36 18.2 72.8 601 4 US-09-949-016-61079 Sequence 37 18.2 72.8 723 4 US-09-016-134-1422 Sequence 38 18.2 72.8 1255 4 US-09-949-016-1772 Sequence 39 18.2 72.8 1255 4 US-08-787-091-1 40 18.2 72.8 1346 3 US-08-787-091-7 41 18.2 72.8 8137 4 US-09-566-921-7 41 18.2 72.8 8137 4 US-09-949-016-13514 Sequence 42 17.8 71.2 26 1 US-08-227-370-4 43 17.8 71.2 26 1 US-08-227-370-4 44 17.6 70.4 945 4 US-09-134-000C-186 Sequence 45 17.4 69.6 616 4 US-09-270-767-11434 Sequence | 35 18.6 74.4 3924 4 US-09-762-195-2 Sequence 2, App. 36 18.2 72.8 601 4 US-09-949-016-61079 Sequence 61079, 37 18.2 72.8 1255 4 US-09-949-016-1772 Sequence 1422, 38 18.2 72.8 1255 4 US-09-949-016-1772 Sequence 1, App. 39 18.2 72.8 146 3 US-08-187-091-1 Sequence 1, App. 40 18.2 72.8 8137 4 US-09-566-921-7 Sequence 7, App. 41 18.2 72.8 8137 4 US-09-266-921-7 Sequence 7, App. 42 17.8 71.2 26 1 US-08-227-370-4 Sequence 4, App. 42 17.8 71.2 26 1 US-08-227-370-4 Sequence 4, App. 44 17.6 70.4 945 4 US-09-134-000C-186 Sequence 186, 14 17.4 69.6 616 4 US-09-270-767-11434 Sequence 11434. | 34 18.6 74.4 3924 4 US-09-023-655-1188 Sequence 1168, Sequence 2, April 18.2 72.8 601 4 US-09-949-016-61079 Sequence 61079, 37 18.2 72.8 723 4 US-09-949-016-434-1422 Sequence 61079, 38 18.2 72.8 1255 4 US-09-949-016-1772 Sequence 1772, 39 18.2 72.8 1446 3 US-09-949-016-1772 Sequence 1772, 39 18.2 72.8 1446 3 US-09-949-016-13514 Sequence 1, April 18.2 72.8 8137 4 US-09-949-016-13514 Sequence 1, April 18.2 72.8 8137 4 US-09-949-016-13514 Sequence 13514, 42 17.8 71.2 26 1 US-08-227-3704 Sequence 4, April 18.2 72.8 47698 4 US-09-140-000C-186 Sequence 4, April 18.2 72.8 47698 4 US-09-140-000C-186 Sequence 4, April 18.2 72.8 47698 4 US-09-140-000C-186 Sequence 11434, 42 17.6 70.4 945 4 US-09-270-767-11434 Sequence 11434. | 33 19 76.0 3321 4 US-09-970-966-175 Sequence 115, 4 34 18.6 74.4 3924 4 US-09-023-655-1168 Sequence 1168, 5 35 18.6 74.4 3924 4 US-09-023-655-1168 Sequence 2, App 36 18.2 72.8 601 4 US-09-949-016-61079 Sequence 61079, 7 37 18.2 72.8 72.3 4 US-09-016-434-1422 Sequence 1422, 7 38 18.2 72.8 1255 4 US-09-949-016-1772 Sequence 1772, 7 39 18.2 72.8 1255 4 US-09-787-091-7 40 18.2 72.8 1346 3 US-08-787-091-7 40 18.2 72.8 8137 4 US-09-949-016-13514 Sequence 1772, 7 41 18.2 72.8 8137 4 US-09-949-016-13514 Sequence 13514, 7 42 17.8 71.2 26 1 US-09-949-016-13514 Sequence 4, App 43 17.8 71.2 26 1 US-09-134-000C-186 Sequence 186, 7 45 17.4 69.6 616 4 US-09-270-767-11434 Sequence 11434, 7 45 17.4 69.6 616 4 US-09-270-767-11434 Sequence 11434, 7 47 18.4 18.5 18.5 18.5 18.5 18.5 18.5 18.5 18.5 | 32 19 76.0 3321 4 US-09-825-294-175 Sequence 175, 7 33 19 76.0 3321 4 US-09-970-966-175 Sequence 175, 7 34 18.6 74.4 3924 4 US-09-762-195-2 35 18.6 74.4 3924 4 US-09-762-195-2 36 18.2 72.8 601 4 US-09-762-195-2 37 18.2 72.8 72.3 4 US-09-16-434-1422 Sequence 1422, 7 38 18.2 72.8 1255 4 US-09-16-1772 Sequence 172, 7 39 18.2 72.8 1255 4 US-09-949-016-1772 Sequence 1772, 7 40 18.2 72.8 8137 4 US-09-787-195-1 Sequence 7, Apple 18.2 72.8 8137 4 US-09-949-016-13514 Sequence 7, Apple 18.2 72.8 8137 4 US-09-949-016-13514 Sequence 7, Apple 18.2 72.8 8137 4 US-09-26-21-7 Sequence 7, Apple 18.2 72.8 8137 4 US-09-26-21-7 Sequence 7, Apple 18.2 72.8 8137 4 US-09-26-21-3514 Sequence 7, Apple 18.2 72.8 47698 4 US-09-249-016-13514 Sequence 4, Apple 18.2 72.8 47698 4 US-09-27-370-4 Sequence 4, Apple 18.2 72.8 47698 4 US-09-27-370-4 Sequence 4, Apple 18.2 72.8 47698 4 US-09-27-370-4 Sequence 4, Apple 18.2 72.8 47698 4 US-09-27-370-85-1434 Sequence 186, 14.5 17.4 69.6 616 4 US-09-270-767-11434 Sequence 186, 14.5 17.4 69.6 616 4 US-09-270-767-11434 Sequence 11434. | 31 19 76.0 3321 4 US-09-713-590-175 Sequence 175, # 32 19 76.0 3321 4 US-09-825-294-175 Sequence 175, # 33 19 76.0 3321 4 US-09-966-175 Sequence 175, # 34 18.6 74.4 3924 4 US-09-762-195-2 Sequence 1168, # 35 18.6 74.4 3924 4 US-09-762-195-2 Sequence 2, App. 36 18.2 72.8 601 4 US-09-949-016-61079 Sequence 61079, # 37 18.2 72.8 1255 4 US-09-949-011-172 Sequence 1772, # 39 18.2 72.8 1255 4 US-09-949-011-172 Sequence 1772, # 40 18.2 72.8 8137 4 US-09-949-016-13514 Sequence 1, App. 41 18.2 72.8 8137 4 US-09-949-016-13514 Sequence 4, App. 42 17.8 71.2 26 1 US-08-227-370-4 Sequence 4, App. 43 17.6 70.4 945 4 US-09-270-767-11434 Sequence 186, # 45 17.4 69.6 616 4 US-09-270-767-11434 Sequence 11434, # 50 11434, # 50 11434, # 50 1155 Sequence 4, App. 51 11434, # 52 11434, # 53 1144 Sequence 11434, # 54 17.4 69.6 616 4 US-09-270-767-11434 Sequence 11434, # 55 11434, # 56 11434, # | 30 19 76.0 3321 4 US-09-640-173-175 Sequence 175, # 31 19 76.0 3321 4 US-09-625-2175 Sequence 175, # 32 19 76.0 3321 4 US-09-925-224-175 Sequence 175, # 33 19 76.0 3321 4 US-09-925-2175 Sequence 175, # 34 18.6 74.4 3924 4 US-09-923-655-1168 Sequence 1168, # 35 18.6 74.4 3924 4 US-09-923-655-1168 Sequence 2, App. 36 18.2 72.8 601 4 US-09-949-016-61079 Sequence 61079, # 37 18.2 72.8 72.3 4 US-09-949-016-61079 Sequence 1172, * 38 18.2 72.8 1255 4 US-09-949-016-172 Sequence 1172, * 39 18.2 72.8 1255 4 US-09-949-016-172 Sequence 1772, * 39 18.2 72.8 1246 3 US-09-787-091-7 Sequence 1772, * 39 18.2 72.8 1255 4 US-09-949-016-13514 Sequence 1772, * 41 18.2 72.8 8137 4 US-09-949-016-13514 Sequence 13514, * 42 17.8 71.2 26 1 US-09-949-016-13514 Sequence 4, App. 43 17.6 70.4 945 4 US-09-134-000C-186 Sequence 186, * 45 17.4 69.6 616 4 US-09-270-767-11434 Sequence 111434, * 45 17.4 69.6 616 4 US-09-270-767-11434 Sequence 111434, * 45 17.4 69.6 616 4 US-09-270-767-11434 Sequence 111434, * 47 18.4 18.5 18.5 18.5 18.5 18.5 18.5 18.5 18.5 | 29 21.8 87.2 4233 3 US-09-450-105-1 30 119 76.0 3321 4 US-09-713-175 Sequence 175, # 31 19 76.0 3321 4 US-09-713-50-175 Sequence 175, # 32 19 76.0 3321 4 US-09-713-50-175 Sequence 175, # 33 19 76.0 3321 4 US-09-970-966-175 Sequence 175, # 34 18.6 74.4 3924 4 US-09-655-1168 Sequence 175, # 35 18.6 74.4 3924 4 US-09-762-195-2 36 18.2 72.8 601 4 US-09-762-195-2 37 18.2 72.8 72.3 4 US-09-949-0.16-1079 Sequence 1022, 7 38 18.2 72.8 72.3 4 US-09-949-0.16-1772 Sequence 1722, 7 39 18.2 72.8 1255 4 US-09-949-0.16-1772 Sequence 1722, 7 39 18.2 72.8 1346 3 US-08-787-991-1 40 18.2 72.8 8137 4 US-09-949-0.16-13514 Sequence 17, Apple 18.2 72.8 8137 4 US-09-949-0.16-13514 Sequence 1, Apple 18.2 72.8 47698 4 US-09-949-0.16-13514 Sequence 1, Apple 18.2 72.8 47698 4 US-09-349-0.16-13514 Sequence 1, Apple 18.2 72.8 47698 4 US-09-349-0.000-186 Sequence 186, M 45 17.4 69.6 616 4 US-09-270-767-11434 Sequence 186, M |

ALIGNMENTS

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APPLICANT: Stremmel, Wolfgang
TITLE OF INVENTION: Phosphatidylcholine as Medication with
TITLE OF INVENTION: Phosphatidylcholine as Medication with
FILE REFERENCE: 34691/208520
CURRENT APPLICATION NUMBER: US/09/762,195
CURRENT FILING DATE: 2001-02-05
PRIOR APPLICATION NUMBER: PCT/EP99702426
PRIOR APPLICATION NUMBER: PCT/EP99702426
PRIOR APPLICATION NUMBER: 198 35 526 2 DE
PRIOR APPLICATION NUMBER: 198 35 570.8 DE
PRIOR APPLICATION NUMBER: 198 57 570.8 DE
PRIOR SILING DATE: 1998-12-15
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 1
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-762-195-1
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US-08-784-649A-1/c
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Best Local S
Matches 25
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                                                                                                                                                                                                                                                                                                        Sequence 1, Application US/08784649A Patent No. 5830697
                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Sikic, Branimir I
APPLICANT: Chen, Gang
TITLE OF INVENTION: P-GLYCOPROTEIN MUTANT RESISTANT
TITLE OF INVENTION: CYCLOSPORIN MODULATION
NUMBER OF SEQUENCES: 5
  STREET: ZZVV
STREET: ZZVV
CITY: Menlo Park
CTATE: CA
TTATE: TTATE
                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           y Match 100.0%; Score 25; DB 4 Local Similarity 100.0%; Pred. No. 0.01; hes 25; Conservative 0; Mismatches
COUNTRY:
                                                                          ADDRESSEE: Fish & Richardson
STREET: 2200 Sand Hill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                               1553
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COMPUTER READABLE FORM: MEDIUM TYPE: Floppy

Floppy disk

94025

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TELEFAX: 415-00. 5:
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 4264 base pairs
TYPE: nucleic acid
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INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 4264 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
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                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Sherwood, Pamela J
REGISTRATION NUMBER: Reg.No. 5
REFERENCE/DOCKET NUMBER: 06037
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-322-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Sherwood, Pamela J
REGISTRATION NUMBER: ReG.NO. 5830697 36,677
REFERENCE/DOCKET NUMBER: 06037/007001
TELECOMMUNICATION INFORMATION:
TELECHONE: 415-322-5070
                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NERAL INFORMATION:
SIKIC, Branimir I
APPLICANT: Sikic, Branimir I
APPLICANT: Chen, Gang
TITLE OF INVENTION: P-GLYCOPROTEIN MUTANT RESISTANT TO
TITLE OF INVENTION: CYCLOSPORIN MODULATION
                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: 11
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/784,649A
FILING DATE:
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                                                                                                                                        APPLICATION NUMBER:
FILING DATE:
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Menlo Park
STATE: CA
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                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 94025
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   USA
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                                                                                                                                                                                   5830697 36,677
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RESULT 5

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US-08-784-649A-5
                                                                                                                  ; NAME/KEY:
; LOCATION:
US-08-181-471-2
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US-08-181-471-2/c
                                                          Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 2, Application US/08181471 Patent No. 5641508
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                 TELEFAX: 619-792-8477 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/041,553
FILING DATE: 02-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Fitting, Thomas
REGISTRATION NUMBER: 34,163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 4646 base pair
                                                                                                                                                               FEATURE:
                                                                                                                                                                             ANTI-SENSE:
                                                                                                                                                                                           MOLECULE TYPE: HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: AN TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Li, Lin
APPLICANT: Lishko,
                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/0 FILING DATE: 13-JAN-1994 CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear
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                                                                                                                                                                                                                            POPOLOGY:
                                                                                                                                                                                                                                        STRANDEDNESS:
                                                                                                                                                                                                                                                    TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                               TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
1911 GTGACATTTTCACGGCCATAGCGAA 1887
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                 GTGACATTTTCACGGCCATAGCGAA 25
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12526 High Bluff Drive, Suite 300
                                                                                                                                                                                                                                                                    4646 base pairs
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                                                          Conservative
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425..4267
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100.0%; Pred. No.
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METHOD FOR DELIVERING BENEFICIAL
COMPOSITIONS TO HAIR FOLLICLES
                                                                       100.0%;
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                                                         Score 25; DB
Pred. No. 0.0
0; Mismatches
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                                                                                     Length 4646;
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US-09-023-655-1167/c
                                  Sequence 18, Application US/08583276
Patent No. 5837536
GENERAL INFORMATION:
APPLICANT: McDonagh, Kevin T.
APPLICANT: Nienhuie, Arthur
APPLICANT: Tolstoshev, Paul
ITILE OF INVENTION: IMPROVED EXPRE
TITLE OF INVENTION: MULTIDRUG RESI
TITLE OF INVENTION: SELECTION OF C
NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS 6.2
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/023.655
FILING DATE: HERFWYFFF.
CLASCIPTI HERFWYFFF.
                                                                                                                                                                                                                                                    US-08-583-276-18/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1167, Application US/09023655 Patent No. 6607879
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                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: PA
TELECOMMUNICATION: INFORMATION:
TELEPHONE: (650) 855-0555
TELEFAX: (650) 845-4166
INFORMATION FOR SEQ ID NO: 1167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Cocke, Benjamin ...
APPLICANT: Suban G. Stuart
APPLICANT: Jeffrey J. Seilhamer
APPLICANT: Jeffrey J. Seilhamer
APPLICANT: TOPENTION: COMPOSITION FOR THE DETECTION OF BLOOD CELL GENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Zeller, Karen J.
REGISTRATION NUMBER: 37,071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: INCYTE PHARMACEUTICALS, INC.
STREET: 3174 PORTER DRIVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IMMEDIATE SOURCE:
LIBRARY: GENBA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: PALO ALTO
STATE: CALIFORNIA
COUNTRY: USA
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLONE: 9187468
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ENGTH:
                                                                                                                                                                                                                                                                                                                                1911 GTGACATTTTCACGGCCATAGCGAA 1887
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Carella, Byrne, Bain, Gilfillan,
Cecchi & Stewart
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                                                                              IMPROVED EXPRESSION OF HUMAN
MULTIDRUG RESISTANCE GENES AND IMPROVED
SELECTION OF CELLS TRANSDUCED WITH SUCH
                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 25; DB 4; 100.0%; Pred. No. 0.011;
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APPLICATION NUMBER: 08/332,444
FILING DATE: 31-OCT-1994
APPLICATION NUMBER: 07/887,712
FILING DATE: 22-MAY-1992
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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                                                                COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/752,447
FILING DATE: 15-NOV-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: NO. 599408Bnan, Kevin E
REGISTRATION NUMBER: 35,303
REFERENCE/DOCKET NUMBER: 95,1121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: DW4.V2
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/
FILING DATE: 05-JAN-1996
CLASSIFICATION: 435
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 07068
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inc
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                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Roninson, Igor B
TITLE OF INVENTION: Methods and Reagents for Preparing and
TITLE OF INVENTION: Using Immunoligcal Agents Specific for P-glycoprotein
                TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-913-0001
TELEFAX: 312-913-9808
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PS/2
OPERATING SYSTEM: F
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COUNTRY:
                                                                                                                                                                                                                                                                                                                                 CITY: Chicago
STATE: Illinois
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25; Conserva
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ilarity 100.0%;
Conservative 0
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OLECULE TYPE:

CDNA

STRANDEDNESS: single TOPOLOGY: linear

nucleic acid

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RESULT 8
US-09-316-167-1/c
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                                                                                                                                            FILING DATE: 15-NOV-1996
ATTORNEY/AGENT INFORMATION:
NAME: NO. 6365357nan, Kevin E
REGISTRATION NUMBER: 35,303
REFERENCE/DOCKET NUMBER: 95,1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-913-0001
TELEPAX: 312-913-9808
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 4669 base pairs
TYPE: nucleic acid
STEAMNEPHNES: 61716
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Matches
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Patent No. 6365357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: Mechetner, Eugene
APPLICANT: Roninson, Igor B
TITLE OF INVENTION: Methods a
TITLE OF INVENTION: Using Imm
FEATURE:
NAME/KEY:
LOCATION:
                                                                   FEATURE:
NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE
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CORRESPONDENCE ADDRESS
                                                                                                 OLECULE TYPE:
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LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Chicago
STATE: Illinois
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CLASSIFICATION:
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LOCATION: 425..4264
                                                  LOCATION:
                                                                                                                                STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 60606
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                                                                                                                    ropology:
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25; Conservative
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                                                5'UTR
1..424
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4265..4669
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1..424
 CDS
425..4264
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                                                                                                                                                                                                                                                                                                                                                        08/752,447
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                                                                                                                                                                                                                                                                                                       Kevin E
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JUCATION: 4265..4669;
SEQUENCE DESCRIPTION: SEQ ID NO:
US-09-397-233-1
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; LOCATION:
US-09-316-167-1
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                                                               Query Match
Best Local Similarity
Matches 25; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 1, Application US/09397233 Patent No. 6630327 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 100.0%; Score 25; I Best Local Similarity 100.0%; Pred. No. 0. Matches 25; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATION DATA:
SOFTWARE PATENTIN Release #1.0, Vere
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/397,233
FILING DATE: 16-Sep-1999
CLASSIFICATION: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: NO. 6630327nan, Kevin E
REGERENCE/DOCKET NUMBER: 95,1121-C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Mechetner, Eugene
TITLE OF INVENTION: Methods and Reagents for Preparing and
Using Immunological Agents Specific for P-glycoprotein
                                                                                                                                                                                                         FEATURE:
1911 GTGACATTTTCACGCCCATAGCGAA 1887
                                                                                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: 312-913-0001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1911 GTGACATTTTCACGGCCATAGCGAA 1887
                                                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
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                     1 GTGACATTTTCACGGCCATAGCGAA 25
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                                                                                                                                                                                                                                                                          NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                               LENGTH: 4669 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                           NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                            NAME/KEY:
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STATE: Illinois
                                                                                                                                                                                                                          LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 312-913-0002
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                                                                    Conservative
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4265..4669
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                                                                    Mismatches
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                                                                                  DB 4;
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655 Fifteenth Street N.W. Suite 330

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; TITLE OF INVENTION: COMPOSITIONS FOR CLONES CONTAINING DNA
; SEQUENCES ASSOCIATED WITH MULTIDRUG RESISTANCE IN HUMAN CELLS
; NUMBER OF SEQUENCES: 4
; CURRENT APPLICATION DATA:
; CURRENT APPLICATION TOTALE:
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5206352-3/c
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CURRENT APPLICATION DATA:
                                                                                                                                                                                                                            RESULT 12
US-08-793-610-5/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No. 5206352, APPLICANT: Roninson, Igor B., Pastan Ira H., Gottesman,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Michael M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO:3:
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                                                 Sequence 5, Application US/08793610

Patent No. 5858744

GENERAL INFORMATION:
APPLICANT: BAUM, Christopher
APPLICANT: BAUM, Christopher
APPLICANT: OSTERTAG, Wolfram
TITLE OF INVENTION: RETROVIRAL VECTOR HYBRIDS AND THE USE THEROF
TITLE OF INVENTION: FOR GENE TRANSFER
                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/07/622,836
FILING DATE: 24-SEP-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 892,575
FILING DATE: 01-AUG-1986
APPLICATION NUMBER: 845,610
PRIOR DATE: 28-MAR-1986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/07/622,836
FILING DATE: 24-SEP-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 892,575
FILING DATE: 01-AUG-1986
APPLICATION NUMBER: 845,610
FILING DATE: 28-MAR-1986
                                           NUMBER OF SEQUENCES:
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Nikaido, Marmelstein, Murray & Oram LLF
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100.0%; Prf
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Pred. No. 0.011;
Mismatches
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RESULT 13
US-09-306-417-1/c
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)638-5000
TELEPAX: (202)638-4810
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1, Application US/09306417 Patent No. 6548301 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 25; Conserv
                                                                                                                                                                                         SEQ ID NO 1
LENGTH: 8630
                                                                                                                                                                                                                                             FILE REFERENCE: P50491
CURRENT APPLICATION NUMBER: US/09/306,417
CURRENT FILLNG DATE: 1999-05-06
EARLIER APPLICATION NUMBER: DE 198 22 115
EARLIER FILLNG DATE: 1998-05-08
NUMBER OF SEQ ID NOS: 17
                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Retroviral Gene Transfer Vectors
                                                                                                                                                                                                                             SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Heinrich-Pette-Institut
FEATURE:
NAME/KEY: misc feature
LOCATION: (1)...(160)
OTHER INFORMATION: plasmid backbone (pUC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: U.S.A.
ZIP: 20005-5701
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                         OTHER INFORMATION: Description of Artificial Sequence: proviral OTHER INFORMATION: plasmid DNA
                                                                                                                                                     ORGANISM: Artificial Sequence
                                                                                                                                                                           TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: DE 195 03 952.1
FILING DATE: 07-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95/03175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                   FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: DNA
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FILING DATE: 07-MAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: DE P
FILING DATE: 08-SEP-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 655 Fitte
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Berman, Richard J. REGISTRATION NUMBER: 39,105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3303 GTGACATTTTCACGGCCATAGCGAA 3279
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US-09-306-417-2/c
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LENGTH: 8630
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Heinrich-Pette-Institut
TITLE OF INVENTION: Retroviral Gene Transfer Vectors
FILE REFERENCE: P50491
CURRENT APPLICATION NUMBER: US/09/306,417
CURRENT FILING DATE: 1999-05-06
EARLIER APPLICATION NUMBER: DE 198 22 115
EARLIER FILING DATE: 1998-05-08
NUMBER: OF SEQ ID NOS: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc_feature
LOCATION: (1)..(8630)
OTHER INFORMATION: retroviral expression vector SFbeta71m4
                                                                                                                                 NAME/KEY: misc_feature
LOCATION: (161)..(677)
OTHER_INFORMATION: 5'-LTR
                                                                                                                                                                                                        LOCATION: (1)..(160)
OTHER INFORMATION: plasmid backbone (pUC)
                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: misc_feature
LOCATION: (5775)..(8630)
OTHER INFORMATION: plasmid backbone (pUC)
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OTHER INFORMATION: 5'-LTR
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LOCATION: (161)..(677)
NAME/KEY: mat_peptide
LOCATION: (1220)..(5062)
DTHER INFORMATION: mSA1
                                                                                                                                                                                                                                                                                             OTHER INFORMATION: retroviral expression vector SFbeta91mSA1
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OTHER INFORMATION: m4 mdr-1 cDNA
                                                                        JAME/KEY: 5'UTR
JOCATION: (532)..(1219)
                                                                                                                                                                                                                                                                                                     JAME/KEY: misc_feature
JOCATION: (1)..(8630)
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LOCATION: (5215)..(5774)
DTHER INFORMATION: 3'-LTR
                                                                                                                                                                                                                                             AME/KEY: misc_feature
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JOCATION: (532)..(1219)
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les 25; Conserv
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ilarity 100.0%; Pred. No. 0.0
Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                      Description plasmid DNA
    mdr1
    CDNA
                                                                                                                                                                                                                                                                                                                                                                                          of Artificial Sequence: proviral
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Pred. No. 0.012;
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; NAME/KEY: misc_feature; LOCATION: (5775)..(8630); OTHER INFORMATION: plasmid backbone (pUC) US-09-306-417-2
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US-08-793-610-6/c
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               Query Match
Best Local &
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                                                                                                                                                TELEFAX: (202)638-4810
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 9318 base pair
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
NAME/KEY: misc feature
LOCATION: (5215)..(5774)
OTHER INFORMATION: 3'-LTR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)638-5000
                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: PCT/
FILING DATE: 10-AUG-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 08-SEE-17.

PRIOR APPLICATION DATA:

DE 195 03 952.1
                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT AFFLICATION NUMBER: US/US/US/
APPLICATION NUMBER: US/US/US/US/
FILING DATE: 07-MAR-1997
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: DE P 44 31 973.8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Rclease #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 07-FEB-1995 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS: ADDRESSEE: Nikaido,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                   MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
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nes 25; Conserv
                                                                                                     TOPOLOGY: ci
                                                                                                                                  TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: U.S.A.
ZIP: 20005-5701
                                                                                                                                                                                                                                                                      NAME: Berman, Richar REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Washington
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l Similarity
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                                                                                                                                                  9318 base pairs
                                                                                 circular
PE: DNA
                                                                                                                                                                                                   (202) 638-4810
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STOCKING-HARBERS, Carol
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               100.0%; Score 25; DB 2;
100.0%; Pred. No. 0.012;
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Pred. No. 0.012;
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Page 7

₽.8 Search completed: February 9, 2005, 17:11:18 Job time : 79.592 secs

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1: geneseqn1980s:*
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25
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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Match |
| 3860 | 3860 | 3860 | 3860 | 3843 | 3840 | 3840 | 3840 | 3840 | 2307 | 765 | 765 | 765 | 765 | 765 | 765 | 584 | 219 | 25 | 25 | Length |
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| ABA94366 | ABA94365 | AAZ49333 | AAZ49332 | ABQ78185 | ABL91687 | ABX09965 | ABZ35722 | ABV78146 | ABS65230 | ADB92124 | ADB96933 | ADB87950 | ADB20861 | ACF62746 | AAA96071 | ACH69096 | ACH82796 | AAD39012 | AAD39002 | B |
| Aba94366 Human BCK | Human | Human | Human | Human | Human | Abx09965 Human mdr | Human | Human | CUNA | 4 | Human | | Adb20861 MRP1 base | Act62746 Cancer ba | Human E | Ach69096 Human gen | Ach82796 Human gen | Aad39012 Human mdr | Aad39002 Human mdr | Description |

| C 45 | C 44 | | C 42 | | | | c 38 | | | | c 34 | | | c 31 | | c 29 | | | c
26 | | | c 23 | c 22 | c 21 | 2 |
|---------------------|----------|----------|----------|---------------------|----------|--------------------|--------------------|----------|----------|----------|----------|--------------------|----------|----------|----------|--------------------|----------|----------|----------|----------|----------|----------|----------|----------|---|
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| 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | ;
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| 4669 | 4669 | 4669 | 4669 | 4646 | 4646 | 4646 | 4646 | 4646 | 4646 | 4646 | 4646 | 4643 | 4643 | 4643 | 4643 | 4643 | 4643 | 4643 | 4533 | 4378 | 4349 | 4264 | 4264 | 2865 | 2 |
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| ABK52041 | AAV32645 | AAQ52726 | AAN70752 | ADR52847 | ADI31841 | ADK60994 | AAD38994 | ABL68880 | ABL68592 | AAZ94738 | AAQ72872 | ADP54881 | ADO19748 | ADP18689 | ABX77217 | ABV74349 | ABV94267 | ABS76368 | ABS65229 | AAQ04522 | AAH57442 | AAV66534 | AAV66533 | AA2889/3 | *************************************** |
| ADR52041 CDNA effco | - Hullid | | | Adr5284 / Drug ther | | ACK60994 OVALIAN C | AadJ8994 Human mur | | Kidney | | | Adp54881 Human PKO | | Human | | Abv/4349 Human ABC | | CUNA er | | | | | | | Aaz88973 Human MDR |

ALIGNMENTS

AAD39002 standard; DNA; 25

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RRESULT 1
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ID AAD3
XX
AAD3902
AC AAD3
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AC AAD3
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AX
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BH Hypo
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BH Hypo
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CO12
PM W020
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                                                                                                                                                                                                                                                                                                                                                                                                                       Human; haematologic malignancy; multidrug resistance; MDR; SUMO-1; hypoxia inducible factor-1; small ubiquitin-like-modifier; HIF-1; lymphotid disorder; chronic lymphoproliferative disorder; lymphoma; myeloid disorder; lymphocytic leukaemia; thrombocythaemia; myeloma; angiogenic myeloid metaplasia; myeloid leukaemia; gene therapy; polycythaemia vera; hypoxia responsive element; HRE; antisense;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human mdr1-HRE antisense oligonucleotide #5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAD39002; .
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                                                                                                                                     26-OCT-2000; 2000US-0243542P.
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                                                                                                                                                                                                                                                                                                          modified_base
                                                                                                                                                                                                                                                                                                                                                              Synthetic.
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                                                                                                          BGHM ) BRIGHAM & WOMENS HOSPITAL INC.
                                                                                                                                                                                                                                                                                             /*tag= a
                                                                                                                                                                                                                                                                 /mod_base= OTHER
/note= "Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                            backbone; ss.
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Treating a subject (at risk of) having a hematologic malignancy or multidrug resistance, e.g. lymphoma or myeloma, by administering hypoxia

WPI; 2002-471427/50.

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RESULT 2
AAD39012/c
ID AAD390
XX AAD390
XX AAD390
XX Human
XX Human
XX Hymphc
KW Hypoxi
KW Lymphc
KW angiog
KW polyc)
XX Homo (
XX Homo 
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Matches 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; haematologic malignancy; multidrug resistance; MDR; SUMO-1; hypoxia inducible factor-1; small ubiquitin-like-modifier; HIF-1; lymphoid disorder; chronic lymphoproliferative disorder; lymphoma; myeloid disorder; lymphocytic leukaemia; thrombocythaemia; myeloma; angiogenic myeloid metaplasia; myeloid leukaemia; gene therapy; polycythaemia vera; hypoxia responsive element; HRE; ds.
        Treating a subject (at multidrug resistance, e
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                                                                                                                                                                                                              & WOMENS HOSPITAL INC
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hypoxia

expression, cu

Claim 1;

tissues.

SEQ ID NO 15991; 80pp; English

The invention relates to a nucleic acid probe for measuring human gene expression, comprising any of the 27,400 fully defined nucleotide sequences in the specification, or their complements or fragments, and

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RESULT 3
ACH82796
ID ACH8
XX ACH8
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Best Local S
Matches 25
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(RANK/) RANK D I
(HANZ/) HANZEL I
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                                                                                                                                                                                                                                      New human genome-derived single exon nucleic acid probes useful for human gene expression analysis, for identifying or characterizing alternative splicing events, for assessing genomic alterations or as tools for
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Penn
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rank DR,
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Pred. No. 0.013;
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 219
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                                                                                                                                                                                                                                                                      US2003194704-A1
                                                                                                                                                                                    03-APR-2002; 2002US-00029386
                   Penn SG,
                                                        (PENN/) PENN S G.
(RANK/) RANK D R.
(HANZ/) HANZEL D K.
                                                                                                                                          03-APR-2002; 2002US-00029386
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                     Rank DR,
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ilarity 100.0%;
Conservative 0
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                   Hanzel
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Pred. No. 0.02;
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WPI; 2004-119264/12.
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New human genome-derived single exon nucleic acid probes useful for human gene expression analysis, for identifying or characterizing alternative splicing events, for assessing genomic alterations or as tools for surveying tissues.

Claim 15; SEQ ID NO 2291; 80pp; English

The invention relates to a nucleic acid probe for measuring human gene expression, comprising any of the 27,400 fully defined nucleotide encoding at least 8 amino acids of any of the 6888 amino acid sequences (fully defined in the specification, or their complements or fragments, and comprising at least 8 amino acids of any of the 6888 amino acid sequences (fully defined in the specification. The probe is a single exon probe that comprising a plurality of single exon mucleic acid molecule comparison of the probes of the specification of the probes of the specification of the plurality of single exon nucleic acid comprising at least 8 and addressably isolatable or amplifiable from the plurality), a single exon microarray for measuring human gene expression, a vector comprising the single exon microarray for measuring human gene expression, a method of contiguous amino acids of any of the above- mentioned amino acid contiguous amino acids of any of the above- mentioned amino acid contiguous amino acids of any of the above- mentioned amino acid contiguous amino acids of any of the above- mentioned amino acid contiguous amino acids of any of the above- mentioned amino acid contiguous amino acids of any of the above- mentioned amino acid contiguous amino acids of any of the above- mentioned amino acid contiguous amino acids of any subscription, and a computer-readable contiguous amino acid and acids of any subscription, and a computer-readable contiguous amino acid and acids of any subscription, and a computer-readable contains a database having a plurality of records (each record including data on the expression of a single exon microarrays to contiguous the probes are used in identifying and characterising contains their specification, the probes are used in identifying and characterising accordance and probes are used in identifying and characterising specifications in the genomic locus that includes their exon, in assessing allers and characterising contains a contains and characterising and characterising and i

Sequence 584 BP; 166 A; 139 C; 100 G; 179 T; 0 U; 0 Other;

Matches Query Match Local 25; Similarity Conservative 100.0%; Scc 100.0%; Pro ative 0; Score 25; DB 12; Pred. No. 0.024; Mismatches 0 Length 584; Indels 0; Gaps

0

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RESULT 5
AAA96071/c
                                                       AAA96071 standard; DNA; 765 BP
                                       29-JAN-2001 (first entry)
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320

GTGACATTTTCACGGCCATAGCGAA 344

1 GTGACATTTTCACGGCCATAGCGAA 25

Human Pgp ATP binding cassette (ABC) transporter coding sequence. Drug resistance; ATP gradient; chemotheraputic; antibiotic; herbicide; human; ATP binding cassette; ABC transporter;

WO200052144-A1

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RESULT 6
ACF62746/c
ID ACF62746;
XX
ACF62746;
XX
ACF62746;
XX
CACF62746;
XX
CACF62746;
XX
CACCEr based on CYP3A5 rel
XX
CENCEr; CYP3A5; irinotecan
KW Cytochrome p450; subfamily
KW Cytostatic; gene; ds.
XX
CACCER; CYP3A5, irinotecan
KW Cytostatic; gene; ds.
XX
CACCER; CYP3A5, irinotecan
KW Cytostatic; gene; ds.
XX
CACCER; CYP3A5, irinotecan
XX
PD 20-FEB-2003.
XX
PD 20-FEB-2003.
XX
PD 20-FEB-2003.
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PD 20-FEB-2003.
XX
PD 20-FEB-2003; 2002WO-EP0081
XX
PD 23-JUL-2001; 2001EP-00117;
PR 23-JUL-2001; 2001EP-00117;
PR 24-MAY-2002; 2002EP-000117;
PR 23-JUL-2001; 2001EP-00117;
PR 23-JUL-2001; 2001EP-00117;
PR 23-JUL-2001; 2001EP-00117;
PR 24-MAY-2002; 2002EP-000117;
PR 24-MAY-2002; 2002EP-0
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Best Local S
Matches 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cancer; CYP3A5; irinotecan; pharmaceutical; malignant glioma; cytochrome p450; subfamily IIIA; nifedipine oxidase; polypept cytostatic; gene; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          altering the ATP gradient across the biological membrane of the target cell. The method is useful for modulating drug resistance of cells. It useful for increasing the sensitivity of cells to chemotheraputic and antibiotic agents and increasing resistance to herbicides. The present sequence is human Pgp ATP binding cassette (ABC) transporter coding sequence. This sequence was used in the present invention to modulate drug resistance. Note: The present sequence is not shown in the specification, but is referred to via its GenBank accession number
                                                                                                                                                                                                                                            23-JUL-2001; 2001EP-00117608.
24-MAY-2002; 2002EP-00011710.
                                                                                                                                                                                                                                                                                                                       23-JUL-2002; 2002WO-EP008219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cancer based on CYP3A5 related polynucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to a method for increasing or decreasing drug resistance in target bacteria, yeast, plant or mammalian cells by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 15; Page; 85pp; English.
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abrane of target cell.
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        irinotecan for preparation having genome with variant
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ises altering ATP gr
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Pred.
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     of compositions for treating allele comprising cytochrome
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RESULT 7
ADB20861/c
ID ADB20861 standard; DNA; 765 BE
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AC ADB20861;
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AC ADB20861;
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DT 20-NOV-2003 (first entry)
DT 20-NOV-2003 (first entry)
DX MRP1 based cancer related nucl
XX irinotecan; covarian cancer;
KW irinotecan; covarian cancer;
KW variant allele; multidrug resi
KW ds.

OS Unidentified.
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OS Unidentified.
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PM W02003013533-A2.
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PP 23-JUL-2002; 2002WO-EP008200.
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PF 23-JUL-2002; 2002EP-00011710.
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Matches 25
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                                                                                                                                                                                                     Use of irinotecan or its derivative for preparation of a pharmaceutical composition for treating cancer in a subject having a genome with a variant allele comprising a multidrug resistance protein 1 polynucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-JUL-2001; 2001EP-00117608.
24-MAY-2002; 2002EP-00011710.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 irinotecan; colorectal cancer; cervical cancer;
lung cancer; ovarian cancer; pancreatic cancer;
variant allele; multidrug resistance protein 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MRP1 based cancer related nucleic
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Pred. No. 0.026;
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malignant glioma;
MRP1; cytostatic; gene;
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present invention describes a method for the use of irinotecan derivative for the preparation of a pharmaceutical composition ating colorectal, cervical, gastric, lung, ovarian or pancreatic

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                                 The invention relates to the novel use of irinotecan to treat a patient suffering from cancer. This involves determining if the patient has one or more variant alleles of the UGTIA1 gene, and if the patient has one or more of such variant alleles, irinotecan is administered in an increased or decreased amount in comparison to the amount that is administered without regard to the patient's alleles in the UGTIA1 gene. The invention has cytostatic activity. A composition of the invention acts as a topoisomerase I inhibitor. The method is useful for treating a patient, an animal e.g. mouse or a human, preferably African or Asian, suffering from cancer such as colorectal, cervical, gastric cancer, lung, ovarian, pancreatic cancer or malignant glioma. The present sequence is udes in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            irinotecan; cancer; UGT1A1; cytostatic; topoisomerase I inhibitor;
colorectal cancer; cervical cancer; gastric cancer; lung cancer;
ovarian cancer; pancreatic cancer; malignant glioma;
uridine diphosphate glycosyltransferasel member A1; gene; ds.
                                                                                                                                                                                                                                                                                                              Use of irinotecan to treat cancer patient by determining if patient has variant alleles of UGTIA1 gene, administering increased/decreased amounts of irinotecan based on increased/decreased levels of UGTIA1 gene product.
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24-MAY-2002; 2002EP-00011710.
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llarity 100.0%;
Conservative 0
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Pred. No. 0.0
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RESULT 10 ADB92124/c ID ADB921

ADB92124 standard; DNA; 765

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ID ADB969
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                                                                           Query Match
Best Local S
Matches 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a multidrug resistance 1 (MDR1) polynucleotide. A composition of the invention has cytostatic activity. The invention is useful for the preparation of pharmaceutical compositions for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject (preferably human, more preferably African or Asian) or a mouse. The present sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New use of irinotecan for preparation of pharmaceutical compositions for treating cancer in subject having genome with variant allele comprising multidrug resistance 1 polynucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    lung cancer; ovarian cancer; pancreatic camultidrug resistance 1; MDR1; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to the novel use of irinotecan or its derivative for the preparation of pharmaceutical compositions for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 674; 130pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Heinrich
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24-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-JUL-2002; 2002WO-EP008218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO2003013537-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               irinotecan; colorectal cancer; cervical cancer; lung cancer; ovarian cancer; pancreatic cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human MDR1
                                                                                                                                                                                     Sequence 765
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-FEB-2003
                                                                                                                                                                                                                                          invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
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                                                                              25;
                                                                                                       Similarity
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GTGACATTTTCACGGCCATAGCGAA
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                                                                           100.0%;
ilarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry
                                                                                                                                                                               BP; 224 A; 139 C; 182 G; 220 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2001EP-00117608.
2002EP-00011710.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA sequence SEQ ID NO:674.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
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                                                                           Score 25; DE
Pred. No. 0.0
0; Mismatches
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Pred. No.
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                                                                                                                                     DB
                                                                                                               . 026;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 malignant glioma;
Cyp3A5; MRP1; MDR1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
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                                                                                   Indels
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                                                                                   Gaps
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RESULT 11
ABS65230/6
ID ABS65230/
ID ABS65230/
ID ABS65
XX ABS65230/
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                 Human; ss; gene; splice variant; tumour-involved gene; TIG; pharmaceutical composition; cancer; diagnostic; tumour; genendothelial cell; cell differentiation; cell proliferation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates ro a novel use of irinotecan or its derivative for the preparation of a pharmaceutical composition for treating colorectal, cervical, gastric, lung, ovarian or pancreatic cancer, or malignant glioma in a subject having a genome with a variant allele which comprises a multidrug resistance 1 (MDR1) polynucleotide. A composition of the invention has cytostatic activity. The present sequence is used in the
                                                            US2002086384-A1
                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                       cDNA encoding tumour involved gene (TIG) splice variant, NV-31
                                                                                                                                                                                                                                                                                                                                                                                       15-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                ABS65230;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABS65230 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        exemplification
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New use of irinotecan for preparation of pharmaceutical compositions for treating cancer in subject having genome with variant allele comprising multidrug resistance 1 polynucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-342400/32
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24-MAY-2002;
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lung cancer; ovarian cancer; pancreatic cancer; malignant glioma;
multidrug resistance 1; MDR1; cytostatic; human; UGT1A1; MRP1; TOP1; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human MDR1 related DNA sequence SEQ ID NO:674.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-DEC-2003
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                                                                                                                                                                            therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  672
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  l Similarity
25; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     765
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 224 A; 139 C; 182 G;
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2002EP-00011710.
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                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                    composition; cancer; diagnostic; tumour; gene therapy;
ll; cell differentiation; cell proliferation; apoptosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 25; DB 10
100.0%; Pred. No. 0.026;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               674; 104pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2307
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  220 T; 0 U; 0 Other;
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RESULT 12
ABV78146/c
ID ABV781
XX
AC ABV781
XX
DT 15-NOV
XX
DT 15-NOV
XX
KW RNA in

ABV78146 standard;

DNA;

0

15-NOV-2002

(first entry)

Human mdr-1 DNA SEQ ID NO 30

RNA inhibition; dsRNA1; gene expression inhibitor; oncogene; cytostatic;

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cc variants that are all tumour-involved genes (TIGS). The nucleic acids and cc polypeptides are useful for determining the level of a nucleic acids and coplypeptide in a biological sample, for detecting a variant nucleic acid or copypeptide sequence in a biological sample, for determining the level of variant nucleic acid or polypeptide sequence in a biological sample, for determining the level of variant problems in a biological sample, for determining the level of variant problems in a biological sample and for ratio between the level of variant sequence in a composition comprising and the level of the original sequence from which the variant has been varied by alternative splicing in a second composition comprising a carrier and the nucleic acid, is useful for treating diseases (e.g. cancer) that can be ameliorated or cured by concreasing or decreasing the level of the encoded protein. The nucleic acids are also useful for diagnostic purposes, especially for detecting cancer or a predisposition to cancer, for evaluating the state or caggressiveness of cancer disease, in basic research, for understanding composition to cancer, for evaluating the state or composition to cancer, for evaluating the state or composition to cancer stages in targeting or composition to cancer stages in targeting or composition to cancer stages in which cell cycle is non-composition. The polypeptides are useful for identifying compounds capable of binding to the variant product and modulating tip activity and for modulating endothelial differentiation and proliferation, as well as to composition the coding sequences for the new variants (NV) 1-cancer and disclosed
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                                                                                                                                              Matches
                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel nucleic acid sequence, which is an alternative splicing variant tumor involved genes, useful for detecting cancer, predisposition to cancer, for evaluating cancer state and in gene therapy for treating
                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention discloses isolated human nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; Page 60-61;
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16-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13-MAR-2001; 2001US-00805020
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(DAVI/)
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1911
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DB; ABG79700.
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                                                                                        Similarity
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                                  GTGACATTTTCACGGCCATAGCGAA 25
                                                                                                                                              2307
GTGACATTTTCACGGCCATAGCGAA 1887
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                                                                     Conservative
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2000IL-00136154
                                                                                                                                            BP; 662 A; 441 C; 591 G;
                                                                                      100.0%; Score 25;
100.0%; Pred. No.
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                                                                       Mismatches
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                                                                                                                                            612 T; 0
                                                                                        0.032;
                                                                                                          DB 6;
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                                                                                                        Length 2307;
                                                                                                                                            U; 1 Other;
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                                                                     Gaps
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RESULT 13
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                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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26-OCT-2001;
29-NOV-2001;
07-DEC-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to inhibiting expression of a target gene (I) in a cell by introducing an inhibitory RNA (dsRNA) having a double-stranded structure of at most 49 consecutive bases. At least part of one strand (as1) of dsRNA1 is complementary to (I) and at least one end of dsRNA1 has an overhang of 1-4 nucleotides. The method is used to inhibit the expression of a wide range of genes, e.g. oncogenes, cytokine genes etc. in humans, also genes in Plasmodium or in viruses or viroids that are pathogenic for humans, animals or plants. Introducing an overhang into dsRNA greatly increases effectiveness for inhibiting gene expression, both in vivo and in vitro and also increases stability and thus the effective concentration inside the cell. The present sequence is that of
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 3840 BP; 1130 A; 738 C; 957 G; 1015 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Inhibiting expression of target oncogenes, by administering doubt target and having an overhang.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-590671/63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kreutzer R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-JAN-2002; 2002WO-EP000152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-JUL-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                virucide; protozoacide; gene; ds
                                                                                                          Double stranded RNA; dsRNA; RNAi; RNA inhibition; cytostatic; viru protozoacide; gene expression; antisense; tumour; infection; Plasm virus; viroid; anti-GPP; human; HIV; human immunodeficiency virus; Hepatitis C virus; human papilloma virus; gene; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (RIBO-) RIBOPHARMA AG.
                                                                                                                                                                                                             07-FEB-2003
09-JAN-2001; 2001DE-01000588
                                                                                                                                                                                 Human mdr-1 polynucleotide SEQ ID NO
                          18-JUL-2002
                                                     DE10100588-A1
                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                         1484
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     related to the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; 2001DE-01000586.
; 2001DE-01055280.
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; 2001DE-01060151.
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                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 rget gene, useful e.g. for inhibiting double-stranded RNA complementary to
                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                              Score 25;
Pred. No.
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                                                                                                                                                                                  30
                                                                                                                                                                                                                                                                                                                                                                                                                0.036;
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                                                                                                                                                                                                                                                                                                                                                                                                                             6.
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                                                                                                                                         Plasmodium;
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                                                                                                                      Matches
                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Inhibiting expression of target genes, useful by introducing into cells two double-stranded to the target.
                                                                                                                                                                                                                                      Sequence 3840 BP; 1130 A; 738 C; 957 G; 1015 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 13; Page 27-28; 100pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-683450/74.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (RIBO-) RIBOPHARMA
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                                                                                                                      25;
                                                            μ
                                                                                                                                                      Similarity
                                                         GTGACATTTCACGGCCATAGCGAA 25
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                                                                                                                         Conservative
                                                                                                                   100.0%; S
100.0%; E
tive 0;
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Pred. No. 0.036;
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                                                                                                                                                                                Length 3840;
                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           for treating tumors, that are complementary
                                                                                                                         0;
                                                                                                                         Gaps
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1484
GTGACATTTTCACGGCCATAGCGAA 1460
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0

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RESULT 14
ABX09965/c
ID ABX099
XX ABX099
XX ABX099
XX ABX099
XX Oligor
CF 23-JAN
XX Oligor
CF PI OP-JAN
XX DP-JAN
XX OP-JAN
XX CRIBO-
XX CRIBO-
XX CRIBO-
XX Inhibi
PT introce
PT introce
                                                                                                                                                                                       Oligoribonucleotide; interferon; oncogene; cytokine; Id; developmental;
                                                                                                                                            DE10100587-C1.
                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                       Human mdr-1 DNA fragment SEQ ID 30.
                                                                                                                                                                                                                          23-JAN-2003
                                                                                                                                                                                                                                             ABX09965;
                                                                                                                                                                                                                                                             ABX09965
                                                                                                                          21-NOV-2002.
                                                                                                                                                                              inhibition;
                                                                                                                                                                                                                                                             standard;
                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                              human;
                                                                                                                                                                                                                                                             3840
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introduction of complementary do

et genes, e.g. oncogenes, in cells, b double-stranded oligoribonucleotide,

уd

after

WPI; 2002-742209/81.

Kreutzer R, (RIBO-) RIBOPHARMA

Limmer S,

Rost

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Hadwiger

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treating the

09-JAN-2001; 2001DE-01000587. 09-JAN-2001; 2001DE-01000587

Disclosure; Page 32-33; 98pp; German.

target genes, particularly oncogenes, cytokine genes, Id (not defined) protein genes; developmental or prion genes, or genes expressed in pathogenic organisms (particularly plasmodia) or in viruses or viroids (pathogenic in humans, animals or plants). Treating the cells with interferon greatly increases the extent to which dsRNA can inhibit expression of the target genes, and the effect is even greater when dsRNA are modified to increase their stability. ABX09936-ABX10075 represent gene fragments used to illustrate the method of the invention This invention describes a novel method for inhibiting expression of a target gene by introducing into the cell that contains the target gene elast one oligoribonucleotide (daRNAI) that has a double-stranded (da) structure of not more than 49 consecutive nucleotides (nt), where at least a segment of one strand of the ds structure is complementary with the target gene and the cells are treated with interferon before introduction of dsRNAI. The method is used to inhibit expression of at

Sequence 3840 BP; 1130 A; 738 C; 957 G; 1015 T; 0 U; 0 Other;

Query Match Best Local (Matches 1484 ы 25; Similarity GTGACATTTTCACGGCCATAGCGAA 1460 GTGACATTTTCACGGCCATAGCGAA 25 Conservative 100.0%; 0 Score 25; DB 6; Pred. No. 0.036; Mismatches 6 <u>,</u> Length 3840;

0

Gaps

0

유 δ

ВP

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30

cytostatic; virucide; protozoacide; antibacterial; HCV; gene expression; oligoribonucleotide; tumour; pathogen; virus; viroid; cytokine; prion; antisense oligonucleotide; virucide; protozoacide; antibacterial; ds.

ω Hadwiger

Inhibiting gene expression in cells, useful for e.g. treating tumors, introducing double-stranded complementary oligoRNA having unpaired γď

104pp; German.

RESULT 15
ABL91687;
ABL91687;
AC ABL916887;
AC ABL91689 ABL91797) in a complementary with the target;
AC C antisense inhibition of gene exp The invention relates to a method for inhibiting expression of a target gene (ABL91658-ABL91797) in a cell by introducing at least one oligoribonucleotide that has a double-stranded structure consisting of at most 49 sequential nucleotide pairs, with at least part of one strand complementary with the target gene and has at least one end a single-stranded sequent of 1-4 nt. The method provides oligoribonucleotides for attacked is the first of the stranded at the stranded sequent of 1-4 nt. The method provides oligoribonucleotides the stranded is the stranded sequent of 1-4 nt. The method provides oligoribonucleotides the stranded sequent of 1-4 nt. The method provides oligoribonucleotides the stranded is the stranded sequent of 1-4 nt. The method provides oligoribonucleotides the stranded sequent of 1-4 nt.

> ភិនិនិនិនិនិនិ but the oligoribonucleotides may also be directed against genes present in pathogens (e.g. Plasmodium or viruses/viroids, pathogenic on humans, animals or plants) or against cytokine, 1d, developmental or prion genes. The method provides more effective inhibition of gene expression than use of known oligonucleotides, probably because the unpaired overhang increases stability and thus intracellular concentration

Sequence 3840 BP; 1130 A; 738 C; 957 G; 1015 T; 0 U; 0 Other;

Query Match Best Local Matches 25; Similarity Conservative 100.0%; 0, Score 25; DB 6; Pred. No. 0.036; Mismatches 6 0 Length 3840; Indels 0, Gaps

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Search completed: February Job time: 245.69 secs ٥ 2005, 16:05:58

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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seq length: 2000000000
Published Applications NA:*

1: /cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*

2: /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*

3: /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*

4: /cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIE

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| US-10-029-386-15991 US-10-029-386-2291 US-09-805-020-31 US-09-805-020-30 US-09-866-868A-1 US-09-866-866A-3 US-09-805-020-30 US-10-007-2621-2 US-10-007-926A-258 US-09-968-007A-459 | ID |
| Sequence 15991, A Sequence 2291, App Sequence 31, App11 Sequence 30, App11 Sequence 1, App11 Sequence 3, App11 Sequence 3, App11 Sequence 2, App11 Sequence 2, App11 Sequence 2, App11 Sequence 2, App11 Sequence 258, App Sequence 459, App | Description |

ALIGNMENTS

US-10-029-386-15991
Sequence 15991, Application US/10029386
Publication No. US20030194704A1
GENERAL INFORMATION:

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APPLICANT: Penn, Sharron G.

APPLICANT: Hanzel, David K.

APPLICANT: Hanzel, David K.

TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR (
ITITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR (
ITITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR (
ITITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR (
ITITLE OF INVENTION: EXPRESSION ANALYSIS TWO

FILE REFERENCE: AEOMICA-X-2.

CURRENT FILING DATE: 2001-12-20

NUMBER OF SEQ ID NOS: 34288

SOFTWARE: Annomax Sequence Listing Engine vers. 1.1

SEQ ID NO 15991

LENGTH: 219

TYPE: DNA

ORGANISM: Homo sapiens

FEATURE:
OTHER INFORMATION: EXPRESSED IN BADULT LIVER, SIGNAL = 1.1

OTHER INFORMATION: EXPRESSED IN BARROW, SIGNAL = 1.6

OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 1.6

OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 1.6

OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 1.6

OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 1.6

OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 1.6

OTHER INFORMATION: EXPRESSED IN BRAIN SIGNAL = 1.6

OTHER INFORMATION: EXPRESSED
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US-09-805-020-31/c
Sequence 31, Application US/09805020
Publication No. US20020086384A1
GENERAL INFORMATION:
APPLICANT: LEVINE, Zurit
APPLICANT: LEVINE, Zurit
APPLICANT: LEVINE, SPLICE VARIANTS OF ONCOGENES
FILE REFERENCE: 2786-0168P
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                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/805,020
CURRENT FILING DATE: 2001-03-13
NUMBER OF SEQ ID NOS: 72
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Hanzel, David K.

TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR TITLE OF INVENTION: EXPRESSION ANALYSIS TWO FILE REFERENCE: AEOMICA-X-2 CURRENT APPLICATION NUMBER: US/10/029,386

CURRENT FILING DATE: 2001-12-20

NUMBER OF SEQ ID NOS: 34288
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                                                                                                                                                                       LENGTH: 2307

LENGTH: 2307

TYPE: DNA

ORGANISM: Homo sapiens

FEATURE:

PEATURE:

NAME/KEY: misc feature

LOCATION: (1)...(2307)

OTHER INFORMATION: any n = a,c,g,t any unknown or other
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1 GTGACATTTTCACGGCCATAGCGAA 25
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25; Conserv
                                                          In 100.0%; Score 25; DB 9; Length 2307; Similarity 100.0%; Pred. No. 0.025; 25; Conservative 0; Mismatches 0: Trdol ...
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US-10-384-339C-30
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                                                                                                                     US-09-866-866A-1
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PRIOR FILING DATE: 2002-01-09
PRIOR APPLICATION NUMBER: DE 10100586.5
PRIOR FILING DATE: 2001-01-09
PRIOR APPLICATION NUMBER: DE 10155280.7
PRIOR APPLICATION NUMBER: DE 10158411.3
PRIOR APPLICATION NUMBER: DE 10158411.3
PRIOR APPLICATION NUMBER: DE 10158411.3
PRIOR APPLICATION NUMBER: DE 10150151.4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1, Application US/09866866A Patent No. US20020102244A1 GENERAL INFORMATION:
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Best Local Similarity 100.
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                                                                                                                                                                                                SOFTWARE: PatentIn version 3.0 SEQ ID NO 1
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                                        Matches
                                                            Best Local
                                                                          Ouery Match
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PRIOR FILING DATE: 2000-05-31
PRIOR APPLICATION NUMBER: PCT/US99/11825
PRIOR FILING DATE: 1999-05-27
PRIOR APPLICATION NUMBER: 60/086,988
PRIOR FILING DATE: 1998-05-28
NUMBER OF SEQ ID NOS: 27
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CURRENT APPLICATION NUMBER: US/10/384,339C
CURRENT FILING DATE: 2003-03-07
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TITLE OF INVENTION: COMPOSITIONS AND METHODS
FILE REFERENCE: 20200/2002
                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/866,866A CURRENT FILING DATE: 2001-08-30
                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Schuetz, John
TITLE OF INVENTION: A Method of Identifying
FILE REFERENCE: 1340-1-021CIP2
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ORGANISM: Homo sapiens
PUBLICATION INFORMATION
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                                                                                                                                                       TYPE: DNA
                                                                                                                                     ORGANISM: Homo sapiens
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  1 GTGACATTTTCACGGCCATAGCGAA 25
                                                          Similarity
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                                        Conservative
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100.0%; Pred. No.
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Pred. No.
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0.027;
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RESULT 7

US-09-805-020-30/c

; Sequence 30, Application US/09805020

; Publication No. US20020086384A1

; GENERAL INFORMATION:

; APPLICANT: LEVINE, Zurit

TITLE OF INVENTION: SPLICE VARIANTS OF ONCOGENES

; FILE REFERENCE: 2786-0168P

; CURRENT APPLICATION NUMBER: US/09/805,020

; CURRENT FILING DATE: 2001-03-13

; NUMBER OF SEQ ID NOS: 72

; SOFTWARE: Patentin Ver. 2.1

; SEQ ID NO 30

; FURTURE OF SEQ ID NOS: 72
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; Sequence 3, Application US/09866866A
; Patent No. US20020102244A1
; GENERAL INFORMATION:
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, NAME/KEY: misc_feature
; LOCATION: (1). (4533)
; OTHER INFORMATION: any n = a,c,g,t any unknown or other
US-09-805-020-30
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US-10-072-621-2/c

; Sequence 2, Application US/10072621

; Publication No. US20020169137A1
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LENGTH: 3860
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Best Local Similarity
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TITLE OF INVENTION: A Method of Identifying and/or Isolating Stem Cells
FILE REFERENCE: 1340-1-021CIP2
CURRENT APPLICATION NUMBER: US/09/866,866A
CURRENT FILING DATE: 2001-08-30
RRIOR APPLICATION NUMBER: 09/584,586
RRIOR FILING DATE: 2000-05-31
PRIOR PILING DATE: 2000-05-31
PRIOR APPLICATION NUMBER: PCT/US99/11825
PRIOR APPLICATION NUMBER: 60/086,988
PRIOR FILING DATE: 1998-05-29
PRIOR APPLICATION NUMBER: 60/086,988
PRIOR APPLICATION NUMBER: 60/086,988
PRIOR FILING DATE: 1998-05-28
PRIOR FILING DATE: 1998-05-28
PRIOR FILING DATE: 1998-05-28
PRIOR FILING DATE: 1998-05-28
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                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 4533
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Homo sapiens
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Local Similarity 100.0%; Pred. No. 0.028;
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TITLE OF INVENTION: Nucleic Acid Molecules and Proteins For The Identification, TITLE OF INVENTION: Nucleic Acid Molecules and Proteins For The Identification, FILE REFERENCE: WRI-030 CURRENT FILING DATE: 2002-03-14 PRIOR APPLICATION NUMBER: 60/276,025 PRIOR FILING DATE: 2001-03-14 PRIOR APPLICATION NUMBER: 60/325,149 PRIOR APPLICATION NUMBER: 60/325,102 PRIOR APPLICATION NUMBER: 60/326,026 PRIOR APPLICATION NUMBER: 60/324,967 PRIOR APPLICATION NUMBER: 60/311,732 PRIOR APPLICATION NUMBER: 60/311,732 PRIOR APPLICATION NUMBER: 60/311,732 PRIOR APPLICATION NUMBER: 60/325,102 PRIOR APPLICATION NUMBER: 60/325,102 PRIOR APPLICATION NUMBER: 60/325,102 PRIOR APPLICATION NUMBER: 60/325,102 PRIOR APPLICATION NUMBER: 60/323,580 PRIOR APPLICATION NUMBER: 60/325,102 PRIOR APPLICATION NUMBER: 60/323,580 PRIOR APPLICATION NUMBER
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-097-340-1
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APPLICANT: Ronnop, Bruce P.
APPLICANT: Connop, Bruce P.
APPLICANT: Pollard, Michelle
TITLE OF INVENTION: REGULATION OF AMYLOID PRECURSOR PROTEIN EXPRESSION OR A
TITLE OF INVENTION: BY MODIFICATION OF ABC TRANSPORTER EXPRESSION OR A
FILE REFERENCE: 100103.402
CURRENT APPLICATION NUMBER: US/10/072,621
CURRENT FILING DATE: 2002-02-08
NUMBER OF SEQ ID NOS: 10
SOFTWARE: FAStSEQ for Windows Version 4.0
SEQ ID NO 2
TENCHARLES AND ASSESSED TO A SECONDARY AND ASSESSED TO A SECONDARY AS
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Matches 25;
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                                                                                                                                          SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
LENGTH: 4643
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APPLICANT: Manjula GANN
APPLICANT: Sebastian HO
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Rachel E. MEYERS
Michael MORRISEY
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Karen GLATT
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Robert C. BAST, Jr
Karen LU
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Shubhangi KAMATKAR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rosemarie SCHMANDT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Peter VEIBY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ami SEN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 25; DB 13 ilarity 100.0%; Pred. No. 0.028; Conservative 0; Mismatches
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RESULT 11
US-09-968-007A-459/c
; Sequence 459, Application US/09968007A
; Publication No. US20040115625A1
                                                               CURRENT APPLICATION NUMBER: US/09/968,007A
CURRENT FILING DATE: 2001-10-02
PRIOR APPLICATION NUMBER: US/60/237,172
PRIOR FILING DATE: 2000-10-02
PRIOR APPLICATION NUMBER: US/60/237,173
PRIOR APPLICATION NUMBER: US/60/237,278
PRIOR APPLICATION NUMBER: US/60/237,278
PRIOR APPLICATION NUMBER: US/60/237,278
PRIOR FILING DATE: 2000-10-02
PRIOR APPLICATION NUMBER: US/60/237,294
PRIOR APPLICATION NUMBER: US/60/237,295
PRIOR APPLICATION NUMBER: US/60/237,295
PRIOR APPLICATION NUMBER: US/60/237,295
PRIOR APPLICATION NUMBER: US/60/237,295
PRIOR FILING DATE: 2000-10-02
PRIOR FILING DATE: 2000-10-02
PRIOR FILING DATE: 2000-10-02
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CURRENT FILING DATE: 2001-12-07
PRIOR APPLICATION NUMBER: 60/254,090
PRIOR FILING DATE: 2000-12-08
NUMBER OF SEQ ID NOS: 468
SOFTWARE: Patentin Ver: 2.1
SEQ ID NO 258
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Usining Signatifie OF INVENTION: Gene Sets FILE REFERENCE: 689290-71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: GENE EXPRESSION PROFILING OF PRIMARY BREAST CARCINOMAS TITLE OF INVENTION: USING ARRAYS OF CANDIDATE GENES FILE REFERENCE: 1546-R-00
                                                  PRIOR APPLICATION NUMBER: US/60/237,316
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: BERTUCCI, FRANCOIS APPLICANT: HOULGATTE, REMI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 4643
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: OTHER INFORMATION:
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                              FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1908 GTGACATTTTCACGGCCATAGCGAA 1884
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BIRNBAUM, DANIEL
NGUYEN, CATHERINE
VIENS, PATRICE
FERT, VINCENT
SEQ ID NOS: 1001
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b. US20030143539A1
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(mdr/tap), member 1 (ABCB1) gene.
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Pred. No. 0.028;
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Pred. No. 0.028;
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US-10-641-643-1167/c

Sequence 1167, Application US/10641643
Publication No. US20040077003A1
GENERAL INFORMATION:

APPLICANT: Cocks, Benjamin G. Susan G. Stuart

CORRESPONDENCE ADDRESS:
ADDRESSEE: INCYTE PHARMACEUTICALS,
STREET: 3174 PORTER DRIVE

CITY: PALO ALTO STATE: CALIFORNIA COUNTRY: USA ZIP: 94304 NUMBER OF SEQUENCES: 1508

TITLE OF INVENTION: COMPOSITION FOR GENE EXPRESSION

THE DETECTION OF BLOOD CELL

Jeffrey J.

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                                                                                                                                                                ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-968-007A-747
                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/968,007A
CURRENT FILING DATE: 2001-10-02
PRIOR APPLICATION NUMBER: US/60/237,172
PRIOR FILING DATE: 2000-10-02
PRIOR FILING DATE: 2000-10-02
PRIOR FILING DATE: 2000-10-02
PRIOR PELICATION NUMBER: US/60/237,278
PRIOR PELICATION NUMBER: US/60/237,278
PRIOR PILING DATE: 2000-10-02
PRIOR APPLICATION NUMBER: US/60/237,294
PRIOR APPLICATION NUMBER: US/60/237,295
PRIOR APPLICATION NUMBER: US/60/237,316
PRIOR APPLICATION NUMBER: US/60/237,316
PRIOR PILING DATE: 2000-10-02
PRIOR PILING DATE: 2000-10-02
PRIOR APPLICATION NUMBER: US/60/237,316
PRIOR PILING DATE: 2000-10-02
PRIOR PILING DATE: 2000-10-02
PRIOR PILING DATE: 2000-10-02
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US-09-968-007A-747/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Ebner, Reinhard
TITLE OF INVENTION: Cancer Gene Determination
TITLE OF INVENTION: Gene Sets
FILE REFERENCE: 689290-71
                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 1001
SOFTWARE: PatentIn version
SEQ ID NO 747
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LENGTH: 464
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Publication No.
                                                                                  Matches
                                                                                                                         Query Match
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Best Local Similarity
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                                                                                Local Similarity
es 25; Conserv
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1911 GTGACATTTTCACGGCCATAGCGAA 1887
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                                         1 GTGACATTTTCACGGCCATAGCGAA 25
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o. US20040115625A1
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                                                                                  Conservative
                                                                                                 100.0%; Score 25;
100.0%; Pred. No.
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                                                                                  Mismatches
                                                                                                     0.028;
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                                                                                                                         Length 4646;
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PRIOR APPLICATION NUMBER: 60/222,313
PRIOR FILING DATE: 2000-08-01
NUMBER OF SEQ ID NOS: 18
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1
SEQ ID NO 1
SEQ ID NO 1
TENGTH: 4646
TYPE: DNA
NORANISM: Homo Gapiens
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US-10-343-657-1/c
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                                                                                                                                                                            ; FEATURE:
; NAME/KEY: CDS
; LOCATION: (425)..(4264)
US-10-343-657-1
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FILING DATE: <Unknown>
ATTORNEY AGENT INFORMATION:
NAME: Zeller, Karen J.
REGISTRATION NUMBER: 37,071
REGISTRATION NUMBER: PA-0001 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 855-0555
TELEPAX: (650) 845-4166
INFORMATION FOR SEQ ID NO: 1167:
SEQUENCE CHARACTERISTICS:
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Best Local Similarity 100.
Matches 25; Conservative
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Publication No. US20040086882A1
                                                                                                         Query Match
Best Local S
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APPLICANT: Ruth, Adam
TITLE OF INVENTION: Mutations of the MDR1 P-glycoprotein that Improve Its
TITLE OF INVENTION: Ability to Confer Resistance to Chemotherapeutic Drugs
FILE REFERENCE: 00,616-A
CURRENT APPLICATION NUMBER: US/10/343,657
CURRENT FILING DATE: 2003-10-17
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS 6.2
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/641,643
FILING DATE: 14-Aug-2003
FILING DATE: 14-Aug-2003
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
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SEQUENCE DESCRIPTION: SEQ ID NO: 1167:
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  1911 GTGACATTTTCACGGCCATAGCGAA 1887
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                                       1 GTGACATTTTCACGGCCATAGCGAA 25
                                                                                 ch 100.0%; Score 25; DB 17; Length 4646; 1 Similarity 100.0%; Pred. No. 0.028; 25; Conservative 0; Mismatches 0; Indels 0
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TYPE: nucleic acid
STRANDEDNESS: single
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APPLICANT: Burczynski, Michael
APPLICANT: Twine, Natalie
APPLICANT: Treine, Andrew
APPLICANT: Trejcchio, William
TITLE OF INVENTION: Method for Monitoring Drug Activities In Vivo
FILE REFERENCE: AM101080 (031896-013000)
CURRENT APPLICATION NUMBER: US/10/775,169
CURRENT FILING DATE: 2004-02-11
NUMBER OF SEQ ID NOS: 5278
SOFTWARE: PatentIn version 3.2
SEQ ID NO 198
LENGTH. Acc.
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US-10-775-169-198/c
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Publication No. US20040175743A1
GENERAL INFORMATION:
                                                                                                                                 Best Local Similarity
Matches 25; Conserv
                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                 LENGTH: 4646
TYPE: DNA
ORGANISM: Homo sapiens
                                                      1911 GTGACATTTTCACGGCCATAGCGAA 1887
                                                                                             1 GTGACATTTTCACGGCCATAGCGAA 25
                                                                                                                             100.0%; Score 25; DB 18; ilarity 100.0%; Pred. No. 0.028; Conservative 0; Mismatches 0:
9, 2005, 22:26:40
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Search completed: February Job time : 267.667 secs

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Listing first 45 summaries
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Match Length DB
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gb est2:
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| AU149941 | BQ028054 | AV648804 | AQ194770 | BM684114 | BM719759 | AQ194986 | AI921934 | AV695293 | BM856196 | AI524590 | BI409869 | BI661427 | BE913573 | BI181199 | BI660209 | BG915267 | CK460130 | CK460832 | B1092416 | CN159932 | |
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ALIGNMENTS

RESULT 1 BP220580/c LOCUS SOURCE ORGANISM ACCESSION VERSION KEYWORDS JOURNAL COMMENT REFERENCE DEFINITION 밁 Ś ORIGIN FEATURES TITLE Query Match Best Local S Matches AUTHORS source Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; 8 1 (bases 1 to 570) 8 Suzuki,Y., Yamashita,R., Shirota,M., Sakakibara,Y., Chiba,J., Mizushima-Sugano,J., Nakai,K. and Sugano,S. Sequence comparison of human and mouse genes reveals a homologous block structure in the promoter regions Genome Res. 14 (9), 1711-1718 (2004) Contact: Yutaka Suzuki Department of Virology Institute of Medical Science, University of Tokyo 4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan Email: ysuzuki@ins.u-tokyo.ac.jp. 25; 92 BP220580 BP220580 Sugano cDNA library, colon Homo sapiens cDNA clone COL03395, mRNA sequence. BP220580.1 GI:52093485 EST. 1 GTGACATTTTCACGGCCATAGCGAA 25 Homo sapiens Homo sapiens (human) Similarity GTGACATTTTCACGGCCATAGCGAA Conservative /organism="Homo sapiens" /mol_type="mRNA" /db_xref="taxon:9606" /clone="COL03695" /tissue_type="colon" /clone_Tib="Sugano cDNA library, colon" Location/Qualifiers 100.0%; Score 25; DB 5 100.0%; Pred. No. 0.25; tive 0; Mismatches 83 5 0 Length 570; Indels <u>,</u> Gaps 0;

RESULT 2 AY408954/c

LOCUS DEFINITION

AY408954

Homo sapiens ABCB1 gene,
genomic survey sequence.

3843 bp , VIRTUAL

DNA linear G TRANSCRIPT, partial

GSS 15-DEC-2003

sequence,

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                                                                                                                                                                                                  Clark, A.G., Glanowski, S., Nielson, R., Thomas, I Todd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F Ferriera, S., Wang, G., Zheng, X.H., White, T.J., Adams, M.D. and Cargill, M.
Inferring nonneutral evolution from human-chim
                   Clark,A.G., Glanowski,S., Nielson,R., Thomas,P., Kejariwal,A., Todd,M.A., Tanenbaum,D.M., Civello,D.R., Lu,F., Murphy,B., Ferriera,S., Wang,G., Zheng,X.H., White,T.J., Sninsky,J.J., Adams,M.D. and Cargill,M.
Direct Submission
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Clark, A.G., Glanowski, S., Nielson, R., Thomas, P., Kejariwal, Todd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F., Murphy, B., Ferriera, S., Wang, G., Zheng, X.H., White, T.J., Sninsky, J.J., Adams, M.D. and Cargill, M.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                            gene trios
Science 302 (5652), 1960-1963 (2003)
                                                                                                                                                                                                                                                                                                                                                                     Pan troglodytes (chimpanzee)
Pan troglodytes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted (16-NOV-2003) Celera Genomics, 45 West Gude Drive, Rockville, MD 20850, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Clark, A.G., Glanowski,S., Nielson,R., Thomas,P., Kejariwal,A., Todd,M.A., Tanenbaum,D.M., Civello,D.R., Lu,F., Murphy,B., Ferriera,S., Wang,G., Zheng,X.H., White,T.J., Sninsky,J.J., Adams,M.D. and Cargill,M.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Pan.
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AY408954.1 GI:39764922
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based on alignment.
Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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(16-NOV-2003) Celera Genomics,
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/locus_tag="HCM3396"
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100.0%; Pred. No. 0.37;
Live 0; Mismatches
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                                                                                                                                                                                                    human-chimp-mouse orthologous
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                                                                                                                                                                                                                                          Thomas, P., Kejariwal, A., R., Lu, F., Murphy, B., te, T.J., Sninsky, J.J.,
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This sequence was made by sequencing genomic exons
them based on alignment.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                  CONA Library Preparation: Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLCM2492 row: c column: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 (bases 1 to 929)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
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AGENCOURT 8627902 NIH MGC 43
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Location/Qualifiers
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                                                                                                                    /tissue_type="normal pigmented retinal epithelium"
/lab host="DH10B (phage-resistant)"
/clone_1lb="NIH_MGC_43"
/clone_1lb="NIH_MGC_43"
/clone_1lb="NIH_MGC_43"
/clone="Organ: eye; Vector: pOTB7; Site_1: XhoI; Site_2: EcoRI; cDNA made by oligo-dT priming. Directionally cloned into EcoRI/XhoI sites using the following 5'
adaptor: GGCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library. | "
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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Sus scrofa
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Plate: 00278 row: b column: 7.
Location/Qualifiers
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Amgen EST Program
Amgen Rat EST Program
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CB727009
                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalla; Eutheria; Cetartiodactyla; Suina; Suidae; 1 (bases 1 to 784)
Caetano, A.R., Johnson, R.K. and Pomp, D.
Generation and sequence characterization of a normal library from swine ovarian follicles
Mamm. Genome 14 (1), 65-70 (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BI184814 784 bp mRNA lir
UNL-P-FN-cr-g-09-0-UNL.81 UNL-P-FN Sus scrofa
UNL-P-FN-cr-g-09-0-UNL 3', mRNA sequence.
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BI184814.1 GI:14659223
                                                      Contact: Pomp, D
Department of Animal Sci
University of Nebraska,
Lincoln, NE 68583-0908,
Tel: 402 472 6416
Fax: 402 472 6362
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/clone lib="W Rat hypothalamus (10471)"
/note="Vector: pSPORT1; Site_1: Sal1; Site_2: Not1;
/note="Vector: pSPORT1; Site_1: Sal1; Site_2: Not1;
/notea="Vector: pSPORT1; Site_2: Not2; Site_2: Not2
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/mol_type="mRNA"
/db_xref="taxon:10116"
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Seg primer: M13 -29
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
1 (bases 1 to 894)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collecting of Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
Email: cgapbs-remail.nih.gov
Tissue Procurement: Jeffrey E. Green, M.D.
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                                                                                                                                                                                                        found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov plate: LLAN9798 row: 1 column: 02
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/db xerf="taxon:9823"

/dev_stage="ADULT"

/lab host="DH10B (Life Technologies)"

/lab host="DH10B (Life Technologies)"

/lab host="Wector: pT7T3D-Pac (Pharmacia) with a modified / note="Vector: pT7T3D-Pac (Pharmacia) with gorcine library is a normalized library representing porcine ovarian follicles, ranging between 2.0 to 10.0 mm in ovarian follicles, ranging between 2.0 to 10.0 mm in ovarian follicles, ranging between 2.0 to 10.0 mm in the library will-p-F2. The tag is a string of 5-6 mucleotides present between the Not I site and the oligo-dT track. The library was constructed as described on parallo, Lennon and Soares, Genome Research 6:
                                                                                                                                                               quality sequence stop: 651.
Location/Qualifiers
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/mol_type="mRNA"
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/clone="IMAGE:4218385"
/lab_host="DH108 (T1 phage-resistant)"
                                                db_xref="taxon:10090"
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|strain="FVB/N"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Carninci,P., Shibata,Y., Hayatsu,N., Sugahara,Y., Shibata,K., Itoh,M., Konno,H., Okazaki,Y., Muramatsu,M. and Hayashizaki,Y. Normalization and subtraction of cap-trapper-selected cDNAs to prepare full-length cDNA libraries for rapid discovery of new genome Res. 10 (10), 1617-1630 (2000)
Adachi,J., Aizawa,K., Akimura,T., Arakawa,T., Bono,H., Carni Fukuda,S., Furuno,M., Hanagaki,T., Hara,A., Hashizume,W., Hayashida,K., Hayatsu,N., Hiramoto,K., Hiraoka,T., Hirozane, Hori,F., Imotani,K., Ishii,Y., Itoh,M., Kagawa,I., Kasukawa,
                                                                                     Group Phase I & II Team.
Analysis of the mouse transcriptome of 60,770 full-length cDNAs
Nature 420, 563-573 (2002)
6 (bases 1 to 3182)
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High-efficiency full-length cDNA cloning
Meth. Enzymol. 303, 19-44 (1999)
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/note="Organ: Colon; Vector: pcWV-SPORT6; Site_1: NotI;
Site_2: Sall; cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.6 kb. Constructed by Life
Technologies. Note: this is a NCI_CGAP Library."
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Direct Submission

Submitted (16-JUL-2001) Yoshihide Hayashizaki, The Institute of Submitted (16-JUL-2001) Yoshihide Hayashizaki, The Institute of Submitted (16-JUL-2001) Yoshihide Hayashizaki, The Institute Physical and Chemical Research (FIKEN Genomic Sciences Canter (1806)

Exploration Research Group, RIKEN Genomic Sciences Canter (1807)

RIKEN Yokohama Institute; 1-7-22 Suehiro-cho, Tsurumi-ku, Yoko Kanagawa 230-0045, Japan (B-mail:genome-res@gsc.riken.jp, Kanagawa 230-0045, Japan (B-mail:genome-res@gsc.riken.
                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleo. Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; (bases 1 to 3816)
Clark, A.G., Glanowski, S., Nielson, R., Thomas, P., Kejariwal, J. Todd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F., Murphy, B., Ferriera, S., Wang, G., Zheng, X.H., White, T.J., Sninsky, J.J., Adams, M.D. and Cargill, M.
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Mus musculus ABCB1 gene,
                                                                          Science 302
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                                                                                                                                           Inferring nonneutral evolution from human-chimp-mouse orthologous
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Mus musculus
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URL:http://fantom.gsc.riken.jp/.
Location/Qualifiers
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Please visit our web site for further details.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cDNA library was prepared and sequenced in Mouse Genome
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Fax:81-45-503-9216)
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member 1B (MGD|MGI:97568, GB|NM_011075,
99%, match=2587)"
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/mol_type="mRNA"
/strain="C57BL/6J"
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/clone_Tib="RIKEN full-length er
/dev_stage="11 days pregnant ad
i. .3182
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/db_xref="taxon:10090"
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Direct Submission
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genomic survey sequence.
CL388323.1 GI:51440283
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Contact: Lawrence B. Schook
Department of Animal Sciences
University of Illinois at Urbana Champaign
1201 W. Gregory Dr., Urbana, IL 61801, USA
Tel: 217 265 5326
Fax: 217 244 5617
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                                                                                                                                                                                                   Clones are derived from the porcine BAC library RPCI-44 (http://www.bacpac.chori.org/porcine242.htm). For BAC library availability, please contact Pietter de Jong (pdejon@chori.org). Clones may be purchased from BACPAC Resources (http://BACPACorders.chori.org). This work was undertaken as part of the International Swine Genome Sequencing Consortium by University of Illinois at Urbana Champaign, USA with funds provided by grant No. AG2002-34480-11828 from USDA-CSREES and AG2001-35205-09965 from USDA/NRI (Livestock Genome Sequencing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 411)
Rogatcheva,M.B., Meyers,S., He,W., Larkin,D.M., Marron,B.M.,
Beever,J.E. and Schook,L.B.
Piggy-BACing the Human Genome: Constructing a Porcine Physical Map
Through Comparative Genomics
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Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae;
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                                                                                                                                                                                                                                                                                                                                                                                           Email: schook@uiuc.
                                                                                                                                                                                        Initiative)
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/locue_tag="HCM3396"
/mol_type="genomic DNA"
/strain="four pigs (breed: 37.5% Yorks Landrace and Meishan)"
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mol_type="genomic DNA"
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1 (bases 1 to 674)

1 (bases 1 to 674)

1 noardman, P.E., Sanz-Ezquerro, J., Overton, I.M., Burt, D.W., Bosch, E., Fong, W.T., Tickle, C., Brown, W.R.A., Wilson, S.A. and Hubbard, S.J. A Comprehensive Collection of Chicken cDNAs Curr. Biol. 12 (22), 1965-1969 (2002)
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604145918F1 CSEQRBN11 Gallus gallus
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Gallus gallus
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Department of Biomolecular Sciences
University of Manchester Institute of Science and Technology
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Fax: 01612360409
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PO Box 88, Manchester, M60 1QD, UK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                   /tissue_type="muscle"
//dev stage="adult"
/lab_host="pH10B"
/lab_host="pH10B"
/lab_host="pH10B"
/clone_lib="CSEQRBN11"
/clone_lib="CSEQRBN11"
/clone_lib="CSEQRBN11"
/clone_lib="CSEQRBN11"
/note="Vector: pBluescript_II KS(+); Site_1: ECORI;
/note="Vector: pBluescript_II KS(+); Site_1: PBluescript_II KS(+);
/note="Vector: The library was normalized in 2 rounds using conditions adapted from Soares et al., pNAS (1994) 91: 9228-9232 and Bonaldo et al., Genome Research & (1994) 9: except_that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Simon.Hubbard@umist.ac.uk.
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/note="Vector: pTARBAC2; Site_1: EcoRI; Site_2: EcoRI;
porcine male BAC library produced by Pieter de Jong"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cell_type="blood"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /db_xref="taxon:9823"
/clone="RPCI44_284D11"
a significantly longer reannealing hybridization was
                                                                                                                                                                                                                                                                                                                                                                                                                                    /mol_type="mRNA"
/strain="Layer and broiler"
/db_xref="taxon:9031"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         organism="Gallus gallus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                           sex="Male and female"
                                                                                                                                                                                                                                                                                                                                                                                                                     clone="ChEST986c5"
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cDNA clone ChEST986c5
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Best Local Similarity
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1 (bases 1 to 716)

1 (bases 1 to 716)

1 (bases 1 to 716)

2 (bases 2 to 716)

3 (bases 3 to 716)

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Simon Hubbard
Department of Biomolecular Sciences
University of Manchester Institute
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                                                                    GTGACATTTTCACGGCCATAGCGAA 25
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                                                                                                                                                                                                                                                                                               /clone_lib="CSEQCHN33"
/clone_lib="CSEQCHN33"
/note="Organ: liver; Vector: pBluescript II KS(+); Site_1:
/note="Organ: ligated to NotI adapters, digested with EcoRI Compatible sites of a custom modified MCS of the pBluescript (KS+) vector: The library was normalized in 2 rounds using conditions adapted from Soares et al., PNAS (1994) 91: 9228-9232 and Bonaldo et al., Genome Research 6 (1996): 791, except that a significantly longer reannealing hybridization was used."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Simon. Hubbard@umist.ac.uk.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /mol_type="mRNA"
/strain="Compton Line 15I"
/db_xref="taxon:9031"
/clone="ChEST250k17"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Gallus gallus"
/mol_type="mRNA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /sex="Female"
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Best Local S
Matches 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   545 TGACATTTTCACGGGCATACCGA 523
                                                                                                                                                                         Unpublished (1999)
Contact: Daniela S. Gerhard, Ph.D.
Office of Cancer Genomics
National Cancer Institute / NIH
Bldg. 31 Rm10A07 Bethesda, MD 20892
Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 923)

NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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High quality sequence start: 2
                  CDNA Library Preparation: Invitrogen Corp CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Agencourt Blogetience Corporation Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://inage.lnl.gov plate: NDAM449 row: d column: 08
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AGENCOURT 14205957 NIH MGC 179 Ho
IMAGE:30384175 5', mRNA sequence.
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Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Jag
Email: ysuzukieims_u-tokyo.ac.jp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence comparison of human and mouse genes reveals a homologous block structure in the promoter regions Genome Res. 14 (9), 1711-1718 (2004)
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BP261688 Sugano cDNA library, clone HSI04571, mRNA sequence.
                                                                                                                                                          Tissue Procurement: Dr. Michael Brownstein
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/mol_type="mRNA"
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clone_lib="Sugano cDNA library,
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/clone="HSI04571"
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                        CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM12185 row: f column: 23
High quality sequence stop: 495.
Location/Qualifiers
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1 (bases 1 to 1235)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)

Unpublished (1999)
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  21; Conservative
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/clone_lib="NIH_MGC_179"
/destroyed); Site_2: NotI; Library is oligo-dT primed and directionally cloned (EcoRV site is destroyed upon cloning). Average insert size 1.1 kb. Library was constructed by (Invitrogen). Note: this is a NIH_MGC Library."
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| 108557 Sequence | ARU91275 Sequence | | റ | AX504298 sequence | | | 0 | | Se | σ | ľ | | | | | 2 | 5165 | | | AR399195 Sequence | | | | | 1 0 |

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| RESULT 2 AX504328/c LOCUS DEFINITION ACCESSION VERSION KEYWORDS SOURCE ORGANISM | d
VQ | Query Match
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Matches 2 | source
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FEATURES | REFERENCE
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ORGANISM | RESULT 1 AX504311 LOCUS DEFINITION ACCESSION VERSION |
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| AX504328 Sequence 31 from Patent WO0234291. AX504328.1 GI:23386140 Homo sapiens (human) Homo sapiens | 1 CGAACCAGGGCACGTGCAATGGCGA 25 | 100.0%; Score 25; DB 6; Length 25; Similarity 100.0%; Pred. No. 0.89; 5; Conservative 0; Mismatches 0; Indels 0; | 125
/organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606" | multiple drug resistance Patent: WO 0234291-A 14 02-MAY-2002; THE BRIGHAM AND WOMEN'S HOSPITAL, INC. (US) Location/Qualifiers | Colgan, S.P. Compositions and methods for treating hematologic malignancies | Homo sapiens (human)
Homo sapiens
Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. | AX504311 25 bp DNA linear PAT 27
Sequence 14 from Patent WO0234291.
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| 27-SEP-2002 | | Gaps | | | | Bostom: | 27-SEP-2002 |
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ch 100.0%; Score 25; DB 9; Similarity 100.0%; Pred. No. 0.84; 25; Conservative 0; Mismatches 0
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1 (bases 1 to 215)
Chen,C.J., Clark,D., Ueda,K., Pastan,I., Gottesman,M.M. and Roninson,I.B.
                                                                                                                                                                                                                                                                                                                                                                                              Draft entry and computer-readable sequence by I.B.Roninson, 27-OCT-1989.
                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                            Original source text: Human multidrug resistant cell line
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Genomic organization of the human multidrug resistance (MDR1) gene and origin of P-glycoproteins J. Biol. Chem. 265 (1), 506-514 (1990)
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Human P-glycoprotein (MDR1) gene,
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Patent: WO 0234291-A 31 02-MAY-2002;
THE BRIGHAM AND WOMEN'S HOSPITAL, INC. (US)
Location/Qualifiers
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12 of 26
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M29433.1 GI:187482
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol type="genomic DNA"
/db_xref="taxon:9606"
/map="7q21"
t <1. .>215
                                                                                 189. .>215
/gene="PGY1"
/note="PGY1, intron
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
                                                                                                                                   note="P-glycoprotein;
number=14
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AX481416/c
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BD171402 3843 bp
Method for predicting side effects
used therefor.
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BD171402.1 GI:28412692
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                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Bhattnagar, S.K., George, A.L. Jr. and Nazarenko, I.
Amplification of nucleic acid sequences
Patent: US 5593840-A 1 14-JAN-1997;
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                                                                                                                                                                                                                                /organism="Homo sapiens"
/mol_type="unassigned DN
/db_xref="taxon:9606"
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/mol_type="unassigned
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AF537133/c
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
1 (bases 1 to 3843)
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AF537133.1 GI:22775295
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Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
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                                                                                                                                                                                                                                                                                                                                                             Direct Submission
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1 (bases 1 to 3852)
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Hanscom, S.R., Xiao, G. and Crespi, C.L.
                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished
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Method for predicting side e
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JP 2002223769-A/1
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1. .3852
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism='Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Homo sapiens"
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                                                                                                                                                                                                                                                                        ocation/Qualifiers
                                                                                                                                                                                                                                                    . .3852
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 25; DB Pred. No. 0.7 % Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .3843
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  effects of immunosuppressant
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6.
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                                                                                                                                                                                                                                                                                                                                   6 Henshaw Street,
                                                           p-glycoprotein'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 3843;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRI 11-SEP-2002
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tranglation="MDLEGDRNGGAEKKNFFKLNNKSKKDKKERKPTVSVFSMFRYSN"

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Macaca fascicularis multidrug resistance p-glycoprotein mRNA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         complete cds.
AF537134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 (bases 1 to 3852)
Stocker, P.J., Steimel-Crespi, D.T., Xiao, G. and Crespi, C.L. Direct Submission
Submitted (09-AUG-2002) BD Biosciences, 6 Henshaw Street,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Macaca fascicularis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AF537134.2 GI:31442762
                                                                                                                                                                                                                                                                                                   Sequence update by submitter on Jun 6, 2003 this sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                           3 (bases 1 to 3852)
Stocker, P.J., Steimel-Crespi, D.T., Xiao, G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Stocker, P.J., Steimel-Crespi, D.T., Xiao, G. and Crespi, C.L. Cynomologous monkey P-glycoprotein mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Macaca fascicularis (crab-eating macaque)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AF537134
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                                                                                                                                                                                                                                                                                                                                                         MA 01801, USA
                                                                                                                                                                                                                                                                                                                                                                                     Submitted (06-JUN-2003) BD Biosciences, 6
                                                                                                                                                                                                                                                                                                                                                                                                                 Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MA 01801, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cercopithecinae; Macaca.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (bases 1 to 3852)
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LTVPFSYLIGAFSVGQAS PSIEAFANARGAAFEIFKIIDNKPSIDSYSKSGHKDNIK
GNLERRNVHFSYPSRKEVKILKGLNIKVOSGOTVALVCHSGCGKSTTVQLMQRLYDPT
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GWQLTLLLAIVPIIA AGVVEMKMLSGQALKDKKELEGAGKIATEAIENFRTVVSLT
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PNTLEGNVTENEVYPFTRLDIPVLGCISLEVKKGGTLALVGSSGCGKSTVVQLLER
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DTVPVMNLEEDMTRYAYYYSGIGAGVLVAAYIQVSFWCLAAGRQIHKIRKQFFHAIMR
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LVILAISPVLGVSAAAWAKILSSFTDKELLAYAKAGAVAEEVLAAIRTVIAFFGQKKE
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ATSALDTESEKVVQEALDKAREGRTCIVIAHRLSTIQNADLIVVFQNGRVKEHGTHQQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FYDPLAGKVLLDGKEIKQLNVQWLRAHLGIVSQEPILFDCSISENIAYGDNSRVVSQE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LLAQKGIYFSMVSVQAGAKRQ"
                                                                                                                                                                                                                                                                         2003 this sequence
Location/Qualifiers
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                                                                                                                 . .3852
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Pred. No. 0.78;
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3852 bp
                                                                                                                                                                                                                                                                                                         version replaced gi:22775297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mRNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 3852;
                                                                                                                                                                                                                                                                                                                                                                                             Henshaw Street, Woburn,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Henshaw Street, Woburn,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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REFERENCE AUTHORS TITLE

JOURNAL

REMARK COMMENT

JOURNAL

FEATURES

SdD

VERSION KEYWORDS

ACCESSION

밁 á

RESULT 8

ocus

ORIGIN

Best Loc Matches

/translation="MDLEGDRNGGAEKKNFFKLNNKSKKDKKERKPTVSVFSMFRYSN WLDKLYMVVGTLAAIIHGAGLPLMMLVFGDMTDTFANAGNLGDLGALLFNNTNSSNIT DTVPVMNLEEDMTRYAYYYSGIGAGVLVAAYIQVSFWCLAAGRQIHKIRKQFFHAIMR

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           VERSION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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                                                                                                                                                                                                                                                                               gene
                                                                                                                                                                                                                                    SgS
                                                                                                                                                                                                                                                                                                                                                                                 source
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 (bases 1 to 3852)
Booth-Genthe, C.L., Roller, C.R.
Cloning of Macaca mulatta MDR1
                                                                                                                                                                                                                                                                                                                                                                                                     Laboratories, WP75A-203, West Point, Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (25-MAR-2004) Drug Metabolism, Merck Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 (bases 1 to 3852)
Booth-Genthe, C.L., R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Macaca mulatta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AY582534.1 GI:46394983
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Macaca mulatta multidrug
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cercopithecinae; Macaca.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Macaca mulatta (rhesus monkey)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AY582534
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EGMYSVDGQDIRTINVRFLREIIGVVSQSEVLFATTIAERIRYGREDVTWDEIEKAVK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FYDPLAGKVLLDGKEIKQLNVQWLRAHLGIVSQEPILFDCSISENIAYGDNSRVVSQE
EIVRAAKEANIHAFIESLPNKYSTRVGDKGTQLSGGQKQRIAIARALVRQPHILLLDE
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/mol_type="mRNA"
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                                                                                                                                                                                                                                                                                                     db_xref="taxon:9544"
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                                                                                                                                                                                                                                      .3852
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            L., Roller, C.R. and Rushmore, T.H.
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REFERENCE

KEYWORDS

VERSION

FEATURES

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Matches

ORIGIN

RESULT 11

SOURCE KEYWORDS VERSION

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

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DEFINITION
ACCESSION
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AX322787/c
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TITLE
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Homo sapiens (human)
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Sequence 1 from Patent WO0192877.
AX322787
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                                                                       AX322789.1 GI:18093767
                                                                                                                                        AX322789
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Method of identifying and/or isolating stem
Patent: WO 0192877-A 1 06-DEC-2001;
ST. JUDE CHILDREN'S RESEARCH HOSPITAL (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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YFKLVTMQTAGNE IELENAADESKSE IDTLENGSHDGGSLIRKRSTRRSVRGSQOD
RKLSTKEALDES I PPVSFWRIMKLNLTEWFYEVVGVFCAI INGGLQPAFAVIFSKI IG
IFTRNDDAETKQQNSNLFSLLFLVLGIVSF ITFFLQGFFFGKAGEILTKKLRYMVFRS
MLRQDVSWFDDPKNTTGALTTRLANDAQVKGAIGSRLAI ITONIANLGTGI ISSLIY
GWQLTLLLLAI VPI IA IAGVVEMMALSGQALKDKKELEGAGKIATEAI ENFRTVVSLIT
GWQLTHLLLAI VPI IA IAGVVEMMALSGQALKDKKELEGAGKIATEAI ENFRTVVSLIT
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GWQLTTLLLLAI VPI IA IAGVVEMMALSGQALKDKKELEGAGKIATEAI ENFRTVVSLIT
GWRETHAVDQSLQVPYRNGLKXAHI FGITTSS FTQAMMYFSYNACFRFGAYLVAHSLM
SFEDVLLVFSAVVFGAMAVGQVSSFAPDYAKAKVSAAHI IMIIEKTPLIDSYSTEGLK
PNTLEGNVTENEVVFNYETKLDIFVLQGLSLEVKKGQTLALVGSSGCGKSTVVQLLER
FYDPLAGKVLLDGKEIKQLMVQMLRAHLGITVSQEP ILFDCSISENIA YGDNSRVVSQE
EI VRAAKEANIHAFIESLPNKYSTRVGDKGTQLSGGQKQRIAIARALVROPHILLLDE
EI VRAAKEANIHAFIESLPNKYSTRVGDKGTQLSGGGKQRIAIARALVROPHILLLDE
ATSALDTESEKVVQBELDKAREGRTCIVIAHRLSTIQNADLIVVFQNGRVKEHGTHQQ
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EGMYSVDGQDIRTINVRFLREIIGVVSQEPVLFATTIAENIRYGREDVTMDEIEKAVK
EANAYDFIMKLPQKFDTLYGERGAQLSGGQKQRIAIARALVRNEKILLLDEATSALDT
                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/mol_type="unassigned DN
/db_xref="taxon:9606"
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LVILAISPILGLSAAAWAKILSSFTDKELLAYAKAGVVAEPULAAIRTVIAFGGOKKE
LERYNKNLEBAKRIGIKKAITANIGIGAAFILIYASYALAFWGTTLVLSKEYSIGOV
LTVFFSVLIGAFSVGQASPBIEAFANARGAAFEIFKIIDNKPSIDSYSKSGHKPDNIK
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                                                                                             3860 bp from Patent WO0192877
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ative 0;
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BD190395/c
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VERSION
KEYWORDS
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BD234199/c
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Wolfgang STREMMEL
OS Homo Sapiens
PN JP 2002522381-A/2
PD 23-JUL-2002
PF 06-AUG-1999 JP 2000563262
PF 06-AUG-1998 DE 198 35 526.2,15-DEC-1998 DI wolfgang stremmel
CC
FH Key Location/Qualifiers.
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Phosphatidylcholine as a m
BD190395
BD190395.1 GI:33000134
JP 2002522381-A/2.
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Method of identifying and/or isolating ste
Patent: WO 0192877-A 3 06-DEC-2001;
ST. JUDE CHILDREN'S RESEARCH HOSPITAL (US)
Location/Qualifiers
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1 (bases 1 to 3924)
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                                        lipid disorders and infl
BD234199
BD234199.1 GI:33043969
JP 2002525111-A/9.
                                                                                      BD234199 3924 bp DNA linear PAT 17-JUL-2003
ATP-binding cassette genes and proteins for diagnosis and remedy of
lipid disorders and inflammatory diseases.
            Homo sapiens (human)
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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06-AUG-1998 DE 198 35 526.2,15-DEC-1998 DE 198 57 750.8 PI
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                                                                                                                                                                                                                                                                                                                                 /organism="Homo sapiens"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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AR380623/c
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Schmitz, G. and Klucken, J.

Schmitz, G. and Klucken, J.

Schmitz, G. and Klucken, J.

ATP-binding cassette genes and proteins for di lipid disorders and inflammatory diseases Patent; JP 2002525111-A, 9 13-AUG-2002;

BAYET AKTIENGESELLSCHAFT

OS Homo sapiens (human)
PN JP 2002525111-A/9
PN JP 2002525111-A/9
PP 21-SEP-1999 UP 2000572359
PF 21-SEP-1999 UP 60/101706
PF 25-SEP-1998 US 60/101706
PG C12N15/09,A61X38/00,A61X45/00,A61X48/00,A

A61P29/00,
PC C12N15/09,A61X38/00,A61X45/00,A61X48/00,A

A61P29/00,A61X38/00,A61X45/00,A61X48/00,A

PC C07X14/47,C07X16/18,C12N1/15,C12N1/19,C12

PC G01N33/53,C12N15/00,C12N5/00,A61X37/02
PC G01N33/53,C12N15/00,C12N5/00,A61X37/02
CC human CDNA of ABCB4 (MDR3)
FH Key

FT SOURCE 1.3924

FT SOURCE 1.3924

FT TOTALLO CONTRACTOR (MDR3)
                                                                                                              1678
AR452557
Sequence 2:
AR452557
                                                                                                                                                                         25;
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AR380623
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                                                                                                                                                                                                                                                                                               1 (bases 1 to 3924)
Cocks, B.G., Stuart, S.G. and Seilhamer, J.J.
Compositions for the detection of blood cell
response gene expression
Patent: US 6607879-A 1168 19-AUG-2003;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                           Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
                                                                                                                                          1 CGAACCAGGCACGTGCAATGGCGA 25
                                                                                                                                                                                                                                                                                                                                                                                               Unclassified.
                                                                                                                                                                                                                                                                                                                                                                                                                              Unknown.
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                                                                                                            CGAACCAGGCACGTGCAATGGCGA 1654
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21-SEP-1999 JP 2000572359
25-SEP-1998 US 60/101706
GERD SCHMITZ, JOCHEN KLUCKEN
C12N15/09, A61K38/00, A61K45/00, A61K48/00, A61P3/06, A61P9/10,
                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1168 from patent US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism='Homo sapiens (human)'.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                  /organism="unknown"
/mol_type="genomic DNA"
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                 from
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100.0%; Pred. No.
                3924 bp
patent US 6677319
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6607879.
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0.78;
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                                                                                                                                                                                                                                                                                                                                                   immunological
                                   PAT 20-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PAT 18-DEC-2003
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VERSION
REYMORDS
SOURCE
Unknown.
ORGANISM
Unknown.
Unclassified.
Winclassified.
REFERENCE 1 (bases 1 to 3924)
AUTHORS
Stremmel, W.
TITLE phosphatidylcholine as medication with protective effect large intestinal mucosa for the string for the strin
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                    Minimum DB seq length: 0
Maximum DB seq length: 200000000
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Perfect score:
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                                                                                                                                                                                                                                                      Database :
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pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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Gapop 10.0 , Gapext 1.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1202784 seqs, 818138359 residues
                                                                                            Issued_Patents_NA:*

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6: /cgn2_6/ptodata/1/ina/backfIles1.seq:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.
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SUMMARIES

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1 | : | No. | 3 | |
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| 20.2 | ۲ | ۲ | ۲ | 21.8 | ۲ | ۲ | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | | Score | | |
| 80.8 | 87.2 | ٠ | 87.2 | 87.2 | 7. | 87. | ٥. | 0 | | 100.0 | | ? | | 100.0 | | .0 | | | 100.0 | ٥. | ٥. | 8 | 8 | 100.0 | 100.0 | 100.0 | | Match | * | P |
| 4233 | 4317 | 4317 | 4279 | 4279 | 4279 | 4279 | 9318 | 8630 | 8630 | 6505 | 4669 | 4669 | 4669 | 4669 | 4669 | 4669 | 4646 | 4646 | 4264 | 4264 | 4195 | æ | 98 | 3924 | 92 | 7 | | Length | | |
| w | 4 | 4. | 4. | 4. | 4. | 4 | N | 4. | 4 | N | თ | σ | 4. | w | N | N | 4 | Н | N | N | 4. | 4 | 4 | 4. | 4 | μ | į | DB | | |
| US-09-120-513-1 | -10-044-67 | -09-672-725C | 9-672-725C-2 | -09-672-725 | -09-672-725C-2 | 50- | 3-610- | 17 | -417- | -08-793-610- | 5206352-3 | 352- | S-09 | -09-316- | 2 | 3-27 | US-09-023-655-1167 | -08-181-4 | P | 4-6 | 9-672-8 | 2-810- | 9-762-195- | US-09-762-195-2 | US-09-023-655-1168 | -08-461-823-1 | | ID | | |
| • | equence 1, | (D | e 26, | Sequence 24, Appl | e 22, | equence 1, Appl | 6 | Ņ | 1, Appi | | | o
O | 9 | Sequence 1, Appli | e
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Ц | e 2, Appl | e 5, Appi | е 1, | e 3, Appi | e 1, Appi | e 1, Appı | Sequence 2, Appli | e 1168, A | equence 1, Appl | | Description | | |

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| 45 | 44 | 43 | 42 | 41 | 40 | 39 | 38 | 37 | 36 | <u>ئ</u> | <u>پ</u> | ω
G | 32 | <u>ω</u> | 30 | 29 | 28 |
| 17.2 | 17.2 | 17.2 | 17.2 | 17.6 | 17.6 | 17.6 | 17.6 | 17.6 | 17.6 | 17.8 | 17.8 | 17.8 | 17.8 | 18.2 | 18.6 | 18.6 | 20.2 |
| 68.8 | 68.8 | 68.8 | 68.8 | 70.4 | 70.4 | 70.4 | 70.4 | 70.4 | 70.4 | 71.2 | 71.2 | 71.2 | 71.2 | 72.8 | 74.4 | 74.4 | 80.8 |
| 17907 | 10690 | 708 | 702 | 6404 | 1341 | 1110 | 813 | 771 | 654 | 121970 | 30847 | 601 | 601 | 690 | 4800 | 4047 | 4233 |
| 4 | W | 4. | 4 | 4 | 4. | 4. | 4 | 4 | 4. | 4. | 44 | 4. | 4. | 4 | N | Ν | ω |
| US-09-949-016-13189 | US-08-961-527-93 | US-09-107-433-1098 | US-09-583-110-92 | US-09-902-540-760 | US-09-902-540-7697 | US-09-328-352-2779 | US-09-252-991A-9462 | US-09-252-991A-9372 | US-09-252-991A-9404 | US-09-949-016-17216 | US-09-949-016-16657 | US-09-949-016-191244 | US-09-949-016-191243 | US-09-540-236-1532 | US-08-612-734B-3 | US-08-612-734B-L | US-09-450-105-1 |
| Sequence 13189, A | | | | | 1 6 9 | | | | | | | | | | | , <u>+</u> | , 1- |

ALIGNMENTS

| STATE: Maryland COUNTY: USA ZIP: 20877 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PATENTIN Release #1.0, Version #1.30 CURRENT APPLICATION NUMBER: US/08/461,823 FILING DATE: 05-JUN-1995 CLASSIFICATION ATA: APPLICATION NUMBER: US 08/168,621 FILING DATE: 16-DEC-1993 PRIOR APPLICATION NUMBER: US 08/168,621 FILING DATE: 16-DEC-1993 PRIOR APPLICATION NUMBER: US 08/168,621 APPLICATION NUMBER: US 08/168,621 FILING DATE: 27-JAN-1993 ATTORNEY/AGENT INFORMATION: NAME: KARTA, Glenn E. NAME: KARTA, Glenn E. RECISTRATION NUMBER: DA-0012 CIP 2 TELECOMMUNICATION INFORMATION: TELEPHONE: 301 208-6997 INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS: LENGTH: 2726 base pairs TELEPAX: 301 208-6997 INFORMATIONESS: double TOPOLOGY: linear MOLECULE TYPE: DNA (genomic) HYPOTHETICAL: NO US-08-461-823-1 | RESULT 1 US-08-461-823-1/c Sequence 1, Application US/08461823 Sequence 1, Application US/08461823 Patent No. 5593840 GENERAL INFORMATION: APPLICANT: Bhatnagar, Satish K. APPLICANT: George Jr., Albert L. APPLICANT: Nazarenko, Irina TITLE OF INVENTION: AMPLIFICATION OF NUCLEIC ACID SEQUENCES NUMBER OF SEQUENCES: 27 CORRESPONDENCE ADDRESS: ADDRESSE: OncorPharm, Inc. STREET: 200 Perry Parkway CITY: Gaithersburg |
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US-09-762-195-2/c
Sequence 2, Application US/09762195
Patent No. 6677319
GENERAL INFORMATION:
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US-09-023-655-1168
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; Sequence 1168, Application US/09023655
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Best Local (
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APPLICANT: Stremmel, Wolfgang
                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (650) 845-41 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                    TOPOLOGY: line
IMMEDIATE SOURCE:
LIBRARY: GENBA
                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Zeller, Karen J.
REGISTRATION NUMBER: 37,071
REFERENCE/DOCKET NUMBER: PA
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS 6.2 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/023,655
FILING DATE: HEREWITH
CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: INCYTE PHARMACEUTICALS, INC.
STREET: 3174 PORTER DRIVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Susan G. Stuart
APPLICANT: Jeffrey J. Seilhamer
TITLE OF INVENTION: COMPOSITION FOR THE DETECTION OF BLOOD CELL GENE
TITLE OF INVENTION: EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Cocks,
APPLICANT: Susan
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: C
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CLASSIFICATION:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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o. 660787
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                                                                                                                                                                                           100.0%; Score 25; DB 4; Length 3924; ilarity 100.0%; Pred. No. 0.089; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
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Pred. No. 0.087;
Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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Sequence 1, Application US/09672810
Patent NO. 6617450
GENERAL INFORMATION:
APPLICANT: STEIMEL-CRESPI, DOROTHY
APPLICANT: CRESPI, CHARLES L.
APPLICANT: CRESPI, CHARLES L.
TITLE OF INVENTION: p-GLYCOPROTEINS
FILE REFERENCE: G0307/7018
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-762-195-2
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US-09-672-810-1/c
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-762-195-1
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. TANGTH: 3924
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CURRENT APPLICATION NUMBER: US/09/762,195
CURRENT FILING DATE: 2001-02-05
PRIOR APPLICATION NUMBER: PCT/EP99702426
PRIOR FILING DATE: 1999-08-06
PRIOR APPLICATION NUMBER: 198 35 526 2 DE
PRIOR FILING DATE: 1998-08-06
PRIOR APPLICATION NUMBER: 198 35 57 570.8 DE
PRIOR APPLICATION NUMBER: 198 57 570.8 DE
PRIOR FILING DATE: 1998-12-15
NUMBER OF SEQ ID NOS: 23
                                                                                                                                                                                                                                                                                                                                                    Query Match 100.0%; Score 25; DB 4; Best Local Similarity 100.0%; Pred. No. 0.089;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 1
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                                                                                                                                                                                                                                                                                                                                 Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/762,195
CURRENT FILING DATE: 2001-02-05
PRIOR APPLICATION NUMBER: PCT/EP99702426
PRIOR FILING DATE: 1999-08-06
PRIOR APPLICATION NUMBER: 198 35 526 2 DE
PRIOR FILING DATE: 1998-08-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 198 57 570.8 DE PRIOR FILING DATE: 1998-12-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Stremmel, Wolfgang
TITLE OF INVENTION: Protective Effect on Large Intestinal Mucosa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Prosphatidylcholine as Medication with TITLE OF INVENTION: Protective Effect on Large Intestinal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILE REFERENCE: 34691/208520
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                                                                                                                                                                                                                                                                                                                                 25;
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                                                                                                                                                                                                                                                                                           1 CGAACCAGGCACGTGCAATGGCGA 25
                                                                                                                                                                                                                                                     CGAACCAGGGCACGTGCAATGGCGA
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                                                                                                                                                                                                                                                                                                                                 Conservative
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Pred. No.
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                                                                                                                                                                                                                                                     1682
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                                                                                                                                                                                                                                                                                                                                                                       DB 4;
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CURRENT APPLICATION NUMBER: US/09/672,810

P-GLYCOPROTEINS AND USES THEREOF

CURRENT FILING DATE: 2000 PRIOR APPLICATION NUMBER:

2000-09-28

US 60/156,921

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; LOCATION: (100)...(3949)
US-09-672-810-3
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PRIOR PILING DATE: 1999-09-28
PRIOR APPLICATION NUMBER: US 60/158,818
PRIOR PILING DATE: 1999-10-12
NUMBER OF SEQ ID NOS: 18
SOFTWARE: Factor 18
                                                                                                                                                      Sequence 1, Application US/08784649A Patent No. 5830697
                                                                                                                                                                                                                                                                                                                                                        Query Match 100.0%; Score 25; DB 4; Best Local Similarity 100.0%; Pred. No. 0.09; Matches 25; Conservative 0; Mismatches 0
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Best Local Similarity 100.0%;
Matches 25; Conservative 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR FILING DATE: 1999-09-28
PRIOR APPLICATION NUMBER: US 60/158,818
PRIOR FILING DATE: 1999-10-12
NUMBER OF SEQ ID NOS: 18
SOFTWARE: FASTSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: G0307/7018
CURRENT APPLICATION NUMBER: US/09/672,810
CURRENT FILING DATE: 2000-09-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: STEIMEL-CRESPI, DOROTHY T.
APPLICANT: CRESPI, CHARLES L.
TITLE OF INVENTION: P-GLYCOPROTEINS AND USES THEREOF
                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: CDS
LOCATION: (100)...(3940)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA ORGANISM: Macaca fascicularis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ENGTH: 4195
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                                             APPLICANT: Sikic, Branimir I
APPLICANT: Chen, Gang
TITLE OF INVENTION: P-GLYCOPROTEIN MUTANT RESISTANT TO
TITLE OF INVENTION: CYCLOSPORIN MODULATION
                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1739 CGAACCAGGGCACGTGCAATGGCGA 1715
                                                                                                                                                                                                                                                                             1748 CGAACCAGGCACGTGCAATGGCGA 1724
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Fish & Richardson
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Pred. No. 0.0
0; Mismatches
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CUANTELLING DATE:
FILING DATE:
FILING DATE:
FILING DATE:
CLASSIFICATION: 435
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Sherwood, Pamela J
REGISTRATION NUMBER: Reg.No. 5830697 36,677
REFERENCE/DOCKET NUMBER: 06037/007001
FILECOMMUNICATION INFORMATION:
TELEPHONE: 415-322-5070
TELEPHONE: 415-854-0875
TELEPHONE: 415-854-0875
TELEPHONE: 415-854-0875
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US-08-784-649A-5/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Sikic, Branimir I
APPLICANT: Chen, Gang
TITLE OF INVENTION: P-GLYCOPROTEIN MUTANT RESISTANT
TITLE OF INVENTION: CYCLOSPORIN MODULATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Sherwood, Pamela J
REGISTRATION NUMBER: Reg.No.
REFERENCE/DOCKET NUMBER: 0603
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-322-5070
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INFORMATION FOR SEQ ID NO:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA;
                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/784,649A
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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CORRESPONDENCE ADDRESS
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STRANDEDNESS: SI
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nes 25; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Menlo Park
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                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
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STREET: 2200 Sand Hill Road
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100.0%; Pred. No. 0.09;
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TOPOLOGY: 11 MOLECULE TYPE:

STRANDEDNESS: single

linear

CDNA

SEQUENCE CHARACTERISTICS:

LENGTH:

i: 4264 base pairs nucleic acid

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US-08-181-471-2
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                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/041

FILING DATE: 02-APR-1993

ATTORNEY/AGENT INFORMATION:

NAME: Fitting, Thomas

REGISTRATION NUMBER: 34,163

REFERENCE/DOCKET NUMBER: ANTO

TELECOMMUNICATION INFORMATION:

TELEPHONE: 619-792-3680

TELEPHONE: 619-792-3680

TELEPAX: 619-792-8477

INFORMATION FOR SEQ ID NO: 2:

SEQUENCE CHARACTERISTICS:

LENGTH: 4646 base pairs
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Best Local S
Matches 25
                                                                    Query Match
Best Local Similarity
                                                      Matches
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                                                                                                                                                                                      MOLECULE TYPE:
HYPOTHETICAL: N
ANTI-SENSE: NO
                                                                                                                                                   FEATURE:
NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Li, Lingna
APPLICANT: Lishko, Valeryi K.
TITLE OF INVENTION: METHOD FOR DELIVERING BENEFICIAL
TITLE OF INVENTION: COMPOSITIONS TO HAIR FOLLICLES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/181,471 FILING DATE: 13-JAN-1994 CLASSIFICATION: 424
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                                                    25;
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1 CGAACCAGGGCACGTGCAATGGCGA 25
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12526 High Bluff Drive, Suite 300
                                                  100.0%; Score 25; DB 1 ilarity 100.0%; Pred. No. 0.09; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                            ANT0029P
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                                                                                  DB 1; Length 4646;
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                                                    Indels
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                                                 Gaps
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; CLONE: G187468
US-09-023-655-1167
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US-09-023-655-1167/c
                                                                                                                                    Sequence 18, Application US/08583276
Patent No. 5837536
                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                         Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (650) 845-4166
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                      GENERAL INFORMATION:
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APPLICANT: Tolstoshev, Paul
TITLE OF INVENTION: MURROVED EXPRESSION OF HUMAN
TITLE OF INVENTION: MULTIDRUG RESISTANCE GENES AND IMPROVED
TITLE OF INVENTION: SELECTION OF CELLS TRANSDUCED WITH SUCH GENES
NUMBER OF SEQUENCES: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS 6.2
CURRENT APPLICATION DATA:
                                                                      APPLICANT:
                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                            IMMEDIATE SOURCE:
LIBRARY: GENBANK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: PATELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Zeller, Karen J.
REGISTRATION NUMBER: 37,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Jeffrey TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Susan G. Stuart
APPLICANT: Jeffrey J. Seilhamer
TITLE OF INVENTION: COMPOSITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                                             Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
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                                                                                                                                                                                                                                                                          1 CGAACCAGGGCACGTGCAATGGCGA 25
                                                                                                                                                                                                                                                                                                         l Similarity
25; Conserv
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CALIFORNIA
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                                                                                  McDonagh, Kevin T.
Nienhuis, Arthur
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Application US/09023655
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                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (650) 855-0555
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Pred. No. 0.09;
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CORRESPONDENCE ADDRESS:

ADDRESSEE:

6 Becker Farm Road

Carella, Byrne, Bain, Gilfillan, Cecchi & Stewart

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US-08-752-447-1/c
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/332,444

PILLING DATE: 31-OCT-1994

APPLICATION NUMBER: 07/887,712

PILING DATE: 22-MAY-1992

INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
LENGTH: 4669 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1, Application US/08752447 Patent No. 5994088
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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FILING DATE: 15-NOV-1996
CLASSIFICATION: 435
ATTORNEY_AGENT INFORMATION:
NAME: NO. 5994088nan, Kevin E
REGISTRATION NUMBER: 35,303
REFERENCE/DOCKET NUMBER: 95,1121
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 6 CITY: ROSeland CITY: New Jersey USA
                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DCS/MS-DOS
SOFTWARE: PATENTIAN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/752,447
FILING DATE: 15-NOV-1996
                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Mechetner, Eugene
APPLICANT: Roninson, Igor B
TITLE OF INVENTION: Methods and Reagents for Preparing
TITLE OF INVENTION: Using Immunoligeal Agents Specific
NUMBER OF SEQUENCES: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/0
FILING DATE: 05-JAN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch diskette
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                                                                                                                                                                                                                                                                                           CITY: Chicago
STATE: Illinois
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                                                                                                                                                                                                                                                                            COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
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                      95,1121
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for P-glycoprotein
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RESULT 13
US-09-316-167-1/c
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Patent No. 6365357
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Mechetner
APPLICANT: Roningon,
                                                                                             TELEPHONE: 312-913-00
TELEFAX: 312-913-9808
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                            CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION UMBER: 08/75:
APPLICATION UMBER: 08/75:
PILING DATE: 15-NOV-1996
ATTORNEY/AGENT INFORMATION:
NAME: NO. 6345357nan, Kev-
REGISTRATION NUMBER: 35,30
                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/316,167
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: cDNA FEATURE:
                                                                                                                                REFERENCE/DOCKET NUMBER: 95
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-913-0001
                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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                                                             SEQUENCE CHARACTERISTICS
LENGTH: 4669 base pai
MOLECULE TYPE: CDNA
               STRANDEDNESS:
TOPOLOGY: lin
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                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
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                                              TYPE: nucleic acid
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DEDNESS: single
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                                                                4669 base pairs
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425..4264
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                                single
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Methods and Reagents for Preparing and
Uging Immunoligcal Agents Specific for P-glycoprotein
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Pred. No. 0.09;
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; FEATURE:
; NAME/KEY:
; LOCATION:
US-09-316-167-1
                                                          NAME/KEY: 3'UTR;
;
LOCATION: 4265..4669
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SEQUENCE DESCRIPTION: SEQ ID NO: 1:
US-09-397-233-1
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US-09-397-233-1/c
US-09-397-233-1/c
; Sequence 1, Application US/09397233
; Patent No. 6630327
; GENERAL INFORMATION:
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Matches
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Best Local Similarity
   Matches
                  Best Local
                                  Query Match
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INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 95,1121-C
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-913-0001
                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: No. 6630327nan, Kevin E
REGISTRATION NUMBER: 35,303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: CUnknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Mechetner, Eugene
Roninson, Igor B
TITLE OF INVENTION: Methods and Reagents for Preparing and
Using Immunological Agents Specific for P-glycoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive
                                                                                                                                     FEATURE:
                                                                                                                                                                                   FEATURE:
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                                                                                                                                                                                                                                    FEATURE:
                                                                                                                                                                                                                                                    MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
   al Similarity
25; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Chicago
STATE: Illinois
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                 LOCATION:
                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                   NAME/KEY:
                                                                                                                                                                                                                  NAME/KEY:
                                                                                                                                                                                                                                                                                                                 LENGTH: 4669 base pairs
                                                                                                                                                                                                    LOCATION:
100.0%; Score 25; DB 4; Length 4669; ilarity 100.0%; Pred. No. 0.09; Conservative 0; Mismatches 0; Indels
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4265..4669
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425..4264
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RESULT 15
5206352-3/c
;Patent No.
Search completed: February 9, 2005, 17:11:19
Job time: 79.592 secs
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TITLE OF INVENTION: COMPOSITIONS FOR CLONES CONTAINING DNA; SEQUENCES ASSOCIATED WITH MULTIDRUG RESISTANCE IN HUMAN CELLS; NUMBER OF SEQUENCES: 4
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO:3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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Best Local
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FILING DATE: 24-SEP-1990
PRIOR APPLICATION DATA:
ADDITION TO THE PRIOR TO T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 892,575
FILING DATE: 01-AUG-1986
APPLICATION NUMBER: 845,610
FILING DATE: 28-MAR-1986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 4669
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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                                                                                                                                                                                                                                                                                                                                                                                     Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                Scoring table:
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                                                                                          N_Geneseq_16Dec04:*
1: geneseqn1980s:*
2: geneseqn1990s:*
3: geneseqn2000s:*
4: geneseqn20101as:*
5: geneseqn2012as:*
6: geneseqn2002as:*
7: geneseqn2002as:*
                                                                                                                                                                                                                                                                                                                                                                                                                 IDENTITY_NUC
Gapop 10.0 , Gapext 1.0
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25
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Copyright (c) 1993 - 2005 Compugen Ltd.
geneseqn2003cs:*
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geneseqn2003bs:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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| _ | 25 | 100.0 | 25 | 6 | AAD39013 | Aad39013 Human mdr |
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| (L) | 25 | 100.0 | 2307 | თ | ABS65230 | Abs65230 cDNA enco |
| • | 25 | 100.0 | 2726 | N | AAQ70907 | Aaq70907 Multidrug |
| un . | 25 | | 2726 | N | AA070916 | |
| o i | 25 | • | 2726 | N | AAT43322 | |
| 7 | 25 | | 3837 | 12 | ADH22602 | Adh22602 cDNA enco |
| æ | 25 | 100.0 | 3840 | σ | ABV78146 | Abv78146 Human mdr |
| ø | 25 | 100.0 | 3840 | σ | ABZ35722 | Abz35722 Human mdr |
| 10 | 25 | | 3840 | თ | ABX09965 | Human |
| 11 | 25 | 100.0 | 3840 | თ | ABL91687 | Human |
| 12 | 25 | 100.0 | 3843 | σ | ABQ78185 | |
| 13 | 25 | 100.0 | 3852 | œ | ABT14496 | Abt14496 Rhesus mo |
| 14 | 25 | 100.0 | 3858 | 12 | ADH22599 | Adh22599 cDNA enco |
| 15 | 25 | 100.0 | 3860 | w | AAZ49332 | Human |
| 16 | 25 | 100.0 | 3860 | w | AAZ49333 | Aaz49333 Human G18 |
| 17 | 25 | 100.0 | 3860 | σ | ABA94365 | Aba94365 Human BCR |
| 18 | 25 | 100.0 | 3860 | o | ABA94366 | Aba94366 Human BCR |
| 19 | 25 | 100.0 | 3924 | w | AAZ94742 | Human |
| 20 | 25 | 100.0 | 3924 | w | AAZ88974 | Aaz88974 Human MDR |
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U | 44 | 43 | 42 | 41 | 40 | 39 | 38 | 37 | 36 | 3
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| 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 |
| 4646 | 4646 | 4646 | 4646 | 4646 | 4643 | 4643 | 4643 | 4643 | 4643 | 4643 | 4643 | 4533 | 4378 | 4349 | 4264 | 4264 | 4195 | 4186 | 4148 | 4043 | 4039 | 3988 | 3924 | 3924 |
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| AAD38994 | ABL68880 | ABL68592 | AAZ94738 | AAQ72872 | ADP54881 | ADO19748 | ADP18689 | ABX77217 | ABV74349 | ABV94267 | ABS76368 | ABS65229 | AAQ04522 | AAH57442 | AAV66534 | AAV66533 | AAF86128 | AAF86127 | ACN41835 | ACN41836 | ACN43502 | AAZ88973 | ADI31842 | ABN95801 |
| Aaqs8994 Human mar | | | | | | Human | Human | | | | | | | | | | | | σ | Humar | ACD43502 Human dia | Aaz889/3 Human Muk | Adi31842 Human CUN | Abn95801 Gene #229 |

ALIGNMENTS

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RESULT 1
AAD39013/c
ID AAD390
Human mdr1 gene HIF-1 binding site DNA #6.
                                                                                                                                       AAD39013 standard;
                                                                                                                          23-SEP-2002
                                                                                                                                AAD39013;
                                                                         misc_binding
                                                                                    Homo sapiens
                                                                                                                          (first entry)
                                                                          Location/Qualifiers
                                                                                                                                       DNA;
                                                                                                                                       25
                                                                                                                                        BP.
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Human; haematologic malignancy; multidrug resistance; MDR; SUMO-1; hypoxia inducible factor-1; small ubiquitin-1!ke-modifier; HIF-1; lymphold disorder; chronic lymphoproliferative disorder; lymphoma; myeloid disorder; lymphocytic leukaemia; thrombocythaemia; myeloma; angiogenic myeloid metaplasia; myeloid leukaemia; gene therapy; polycythaemia vera; hypoxia responsive element; HRE; ds.

02-MAY-2002. 25-OCT-2001; 2001WO-US049856 WO200234291-A2 /bound_moiety= "HIF-1"

(BGHM) BRIGHAM & WOMENS HOSPITAL INC.

26-OCT-2000; 2000US-0243542P.

Colgan SP;

WPI; 2002-471427/50.

Treating a subject (at risk of) having a hematologic malignancy or multidrug resistance, e.g. lymphoma or myeloma, by administering hypoxia inducible factor 1 binding molecules or small ubiquitin-like-modifier-1 binding molecules.

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CC The method involves administering hypoxia inducible factor-1 (HRF-1) binding molecules or small ubiquitin-like-modifier (SUMO)-1 binding complex modulators. mdr1-hypoxia responsive celement (HRE) binding molecules or antisense mucleic acid molecules and SUMO-1 binding molecules or antisense mucleic acid molecules and cc subject having or at risk of developing haematologic malignancy or MDR ce.g. a lymphoid disorder or a mysloid disorder). The lymphoid disorders include lymphocytic leukaemia or chronic lymphoproliferative disorders ce.g. lymphoma, myeloma or chronic hymphoid leukaemia, e.g. angiogenic myeloid metaplasia, essential thrombocythaemia or polycythaemia vera. The invention is used in gene therapy. The present sequence is human mdr1 complex of the myeloid site of the present sequence is human mdr1 cross of the myeloid site of the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
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Treating a subject (at risk of) having a hematologic malignancy or multidrug resistance, e.g. lymphoma or myeloma, by administering hypoxia inducible factor 1 binding molecules or small ubiquitin-like-modifier-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; haematologic malignancy; multidrug resistance; MDR; SUMO-1; hypoxia inducible factor-1; small ubiquitin-1!ke-modifier; HIF-1; lymphoid disorder; chronic lymphoproliferative disorder; lymphoma; myeloid disorder; lymphocytic leukaemia; thrombocythaemia; myeloma; angiogenic myeloid metaplasia; myeloid leukaemia; gene therapy; polycythaemia vera; hypoxia responsive element; HRE; antisense; phosphorothioate backbone; ss.
                                                                                                        WPI; 2002-471427/50
                                                                                                                                                                                                                                                         26-OCT-2000; 2000US-0243542P
                                                                                                                                                                                                                                                                                                        25-OCT-2001; 2001WO-US049856
                                                                                                                                                                                                                                                                                                                                                            02-MAY-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human mdr1-HRE antisense oligonucleotide #6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAD39003 standard; DNA; 25
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                                                                                                                                                                                                        (BGHM ) BRIGHAM & WOMENS HOSPITAL INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
1. .25
                                                                                                                                                                                                                                                                                                                                                                                                                                                  /mod_base= OTHER
/note= "Phosphorothioate backbone"
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Pred. No. 0.084;
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RESULT 3
ABS65230/c
ID ABS652
XX ABS652
XX ABS652
XX ABS652
XX LIFT NOV
DE CDNA e
XX Human;
KW Pharma
KW gene t
XX Gene t
XX US2002
XX US2002
XX Homo s
XX US2002
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Best Local (
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                                                                                    Novel nucleic acid sequence,
tumor involved genes, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; ss; gene; splice variant; tumour-involved gene; TIG; pharmaceutical composition; cancer; diagnostic; tumour; gene therapy; endothelial cell; cell differentiation; cell proliferation; apoptosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a method of treating a subject having or at a of developing a haemacologic malignancy or multidary resistance (MDR). The method involves administering hypoxia inducible factor-1 (HIF-1) binding molecules or small ubiquitin-like-modifier (SUMO)-1 binding molecules or small ubiquitin-like-modifier (SUMO)-1 binding molecules or HIF-1-SUMO-1 complex modulators. mdrl-hypoxia responsive
                                                                                                                                                                                                                 WPI; 2002-635679/68.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cDNA encoding tumour involved gene (TIG) splice variant, NV-31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-NOV-2002
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16-MAY-2000; 2000IL-00136154.
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DAVID A.
ROMANO C.
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25; Conserv
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                                                                                                                                                                                                                                                                            David A,
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                                                        which is an alternative splicing variant for detecting cancer, predisposition to state and in gene therapy for treating
                                                                                                                                                                                                                                                                               Bernstein
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cancer, for cancer.

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Enzymatic amplification of target nucleic acid sequences detect a mutation or allele in the target, e.g. that caus
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RESULT 4
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AC AAQ ફ 뭉 The invention discloses isolated human nucleic acid alternative splicing CC variants that are all tumour-involved genes (TIGs). The nucleic acids and CC polypeptides are useful for determining the level of a nucleic acid or CC polypeptide in a biological sample, for detecting a variant nucleic acid or polypeptide in a biological sample, for detecting a variant nucleic acid or polypeptide sequence in a biological sample, for detecting a variant sequence in a for determining the level of variant sequence in a biological sample and the level of the variant sequence in a first biological sample and the level of the original sequence in a biological sample and for raising antibodies. A pharmaceutical composition comprising a carrier and the nucleic acid, is useful for increasing or decreasing the level of the emoletic acid, is useful for cardes are also useful for diagnostic purposes, especially for detecting cancer or a predisposition to cancer, for evaluating the state or cagraesiveness of cancer disease, in basic research, for understanding the physiological function of the original TIG, in targeting or cayled for the same type of cells which may be helpful for the development company. The polypeptides are useful for identifying compounds capable of binding to the variant product and modulating its activity and for modulate apoptosis either ex vivo or in vivo. The sequences presented in capacity and experiments of the coding sequences for the new variants (NV) 1-36 of the TIGs disclosed Matches Query Match Sequence 2307 Claim 1; Page 60-61; 180pp; English Multidrug-resistance AAQ70907; AAQ70907 standard; Synthetic Multidrug-resistance; MDR-1 gene; 25-MAR-2003 27-MAR-1995 04-AUG-1994 WO9417206-A1 WPI; 1994-264118/32. Bhatnagar SK, 12-NOV-1993; Local 2064 μ. 25; ONCOR Similarity CGAACCAGGGCACGTGCAATGGCGA 25 CGAACCAGGGCACGTGCAATGGCGA Conservative (revised) (first en BP; 662 A; 441 George 93US-00010433 93WO-US010883 DNA; entry) 100.0%; gene MDR-1. Ł 2726 .. ВÞ C; 591 G; Score 25; Pred. No. 0 Mismatches 612 T; 0 U; 1 Other; 0.14; DB 6; Length 2307; 0 Indels 0 Gaps

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 2726 BP; 818 A; 538 C; 642 G;
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                                                                                                               This gene is used as an example of a target DNA in a method for the amplification of nucleic acid sequences in a mixture using various DNA probes/primers. The method may be used to detect a mutation or allele the target, e.g. that that causes a genetic disease. (Updated on 25-MP 2003 to correct PN field.)
                                                                                                                                                                                                                       Enzymatic amplification of target nucleic acid sequences to form exact modified copies - has increased fidelity and can identify point mutatic
                                                                                                                                                                                                                                                                                                                                27-JAN-1993;
16-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                           04-AUG-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Multidrug-resistance gene MDR-1.
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27-MAR-1995
                                                                                                                                                                                       Disclosure; Page 30; 69pp; English.
                                                                                                                                                                                                                                                             WPI; 1994-264122/32.
                                                                                                                                                                                                                                                                                   Bhatnagar SK,
                                                                                                                                                                                                                                                                                                                                                                     25-JAN-1994;
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 144 CGAACCAGGGCACGTGCAATGGCGA 120
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                        CGAACCAGGGCACGTGCAATGGCGA 25
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(first entry)
                                                                                          BP;
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                                                                                                                                                                                                                                                                                   George
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93US-00168621.
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                                                                                            818 A;
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                                                                                            538 C; 642 G;
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Pred. No.
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Pred. No. 0.1
0; Mismatches
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J.14;
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RESULT 7
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                                                                                                                                                                                                                                                                          This sequence represents the multidrug resistance gene (MDR-1) amplified CC by the primers shown in AAT43320 and AAT3321. This sequence can be used CC as a target in the method of the invention, for enzymatically amplifying CC a target nucleic acid (TWA) sequence contained in a nucleic acid or comprises using three primers, one complementary CC polymerase. The method comprises using three primers, one complementary CC of the TWA, which is adjacent to the first primer, and a third which is similar to the first segment of the TWA, and is complementary to a good segment of the TWA, and a fused amplification product is created from the TWA using CC these two primers. The first two primers are hybridised to the TWA, and a fused amplification product is dissociated, and hybridised to the third primer, which is then extended. The extended modified camplification product is hybridised to the first two primers, and these CC amplification product is hybridised to the first two primers, and these CC different label so that the method can be used to detect the presence of a mutation or allele by detecting whether the labelled primer is CC contained within the fused amplification product or the extended camplification product. The method of the invention combines certain CC (PCR), but is improved compared to LCR due to the reduced number of primers needed, and the fact that the entire TWA sequence does not need contained within the fact that the entire TWA sequence does not need to be primers needed, and the fact that the entire TWA sequence does not need to be primers to the sequence of the primers needed.
                                                                                                                                                                          Query Match
Best Local (
                                                                                                                                                             Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Amplification method avoiding strand displacement by polymerase - in the detection of mutation(s) and allele(s) associated with genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-JUN-1995;
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human; multidrug
                                                                                        144
                                                                                                                                                                                                                                                                  known
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                                                                                                            CGAACCAGGGCACGTGCAATGGCGA 25
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standard; cDNA; 3837

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Query Match Best Local Matches

L Similarity 25; Conserv

100.0%; ilarity 100.0%; Conservative

٥, Score Pred.

Mismatches 25; No.

0.14;

DB 12; 0;

Length 3837; Indels

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Gaps

0

CGAACCAGGGCACGTGCAATGGCGA

Sequence

3837

BP; 1130 A; 772 C; 962 G;

973 T; 0 U; 0 Other;

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cc as agonists, antagonists, antibodies, expression vectors and host cells, cin appropriate screening and toxicity assays to assess the effects of cexogenous compounds on TRICH expression. The present invention describes TRICH compositions that are useful in the diagnosts, treatment and prevention of various disorders such as cell proliferative (e.g. arteriosclerosis, cancer), autoimmune/inflammatory (e.g. ADDS, asthma), cc arteriosclerosis, cancer), autoimmune/inflammatory (e.g. Cushing's eyndrome hypothyroidism) and for infections. Accordingly, these TRICH comolecules can be used for gene therapy purposes and exhibit various activities such as cytostatic, antiinflammatory, immunosuppressive. Cantiasthmatic, anticonvulsant, nootropic and neveroprotective. Gurthermore, a microarray is useful in monitoring or measuring protein-protein interactions, drug-target interactions and gene expression treasting. This polynucleotide sequence is a human TRICH cDNA of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention relates to novel isolated polynucleotides identified as human transporters and ion channels (TRICH), and the encoded polypeptides thereof. Specifically, it describes using these TRICH molecules, as well
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-MAY-2002; 2002US-0377435P-
03-MAY-2002; 2002US-0377444P-
05-JUN-2002; 2002US-0386497P-
11-JUN-2002; 2002US-0388180P-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New human transporters and ion channels (TRICH), useful for diagnosing, treating and preventing diseases or conditions associated with the aberrant TRICH expression e.g. cancer, AIDS, atherosclerosis, epilepsy,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Baughn MR, Becha SD, Bullo
Emerling BM, Griffin JA, H
Jin P, Kable AE, Khare R,
Ramkumar J, Richardson TW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         arteriosclerosis; cancer; autoimmune/inflammatory; AIDS; asthma; neurological; epilepsy; stroke; developmental; Cushing's syndrome; hypothyroidism; infection; gene therapy; cytostatic; antiinflammatory; immunosuppressive; antiasthmatic; anticonvulsant; nootropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 12; SEQ ID NO 100; 448pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              P-PSDB; ADH22536.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cDNA encoding a human transporter & ion channel (TRICH) protein SeqID100.
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JA, Hafalia AJA, Ison CH, Jackson AA, Jian
Fee R, Lee S, Mason PM, Marquis JP;
TW, Swarnakar A, Tran UK, Chawla NK;
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RESULT 9
ABZ35722/
ID ABZ3
XX
AC ABZ3
XX
DT 07-F
.XX
DE Hume
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ABV78146/c
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                                                                                                                                                                                                         Query Match
Best Local S
Matches 25
                                                                                                                                                                                                                                                                                                    The invention relates to inhibiting expression of a target gene (I) in a cell by introducing an inhibitory RNA (dsRNA1) having a double-stranded structure of at most 49 consecutive bases. At least part of one strand (asl) of dsRNA1 is complementary to (I) and at least one end of dsRNA1 has an overhang of 1-4 nucleotides. The method is used to inhibit the expression of a wide range of genes, e.g. oncogenes, cytokine genes etc. in humans, also genes in Plasmodium or in viruses or viroids that are pathogenic for humans, animals or plants. Introducing an overhang into dsRNA greatly increases effectiveness for inhibiting gene expression, both in vivo and in vitro and also increases stability and thus the effective concentration inside the cell. The present sequence is that of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Inhibiting expression of target gene, useful e.g. for inhibiting oncogenes, by administering double-stranded RNA complementary to target and having an overhang.
  Human mdr-1
                                                                                                                                                                                                                                                                Sequence 3840 BP; 1130 A; 738 C; 957 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI, 2002-590671/63.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kreutzer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-JAN-2001; 2001DE-01000586.
26-0CT-2001; 2001DE-01055380.
29-NOV-2001; 2001DE-01058411.
07-DEC-2001; 2001DE-01060151.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RNA inhibition; dsRNA1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human mdr-1 DNA SEQ ID NO 30.
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                              07-FEB-2003
                                                                                  ABZ35722 etandard; DNA; 3840
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                                                                                                                                                                                                                                                                                           related to the invention
                                                                                                                                                                                                            25;
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                                                                                                                                                                                                                         Similarity
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                                                                                                                                                                       CGAACCAGGGCACGTGCAATGGCGA 25
                                                                                                                                                     CGAACCAGGCACGTGCAATGGCGA 1613
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 polynucleotide SEQ ID
                                                                                                                                                                                                         100.0%; Score 25; Di
llarity 100.0%; Pred. No. 0.:
Conservative 0; Mismatches
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                              entry)
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gene; ds.
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                                                                                                                                                                                                                                                                  1015
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                                                                                                                                                                                                            0
                                                                                                                                                                                                                                     Length 3840;
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21-NOV-2002 DE10100587-C1

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RESULT 10
ABX09965/c
ID ABX0999
XX ABX099
XX ABX099
XY
DT 23-JAN
XX
DE Human
XX
Oligor
XW prion;
XX
OS Homo E
XX
PN DE101(
XX
PD 21-NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to inhibiting expression of a target gene in a cell CC by introducing at least two oligoribonucleotides (dsRNAI and II), both CC with a double-stranded (ds) structure of at most 49 sequential nucleotide CC pairs. At least part of one strand (Sl, S2) of the ds structures in each CC for dsRNAI and II are complementary to regions in the target gene. The CC method uses antisense inhibition of gene expression using double stranded CC RNA inhibition (RNAi). The method is particularly used to treat tumours CC or infections, especially by Plasmodium or viruses/viroids (pathogenic on CC humans, animals or plants). The method provides more effective inhibition CC of expression than known methods using a single dsRNA, even at very low CC concentrations. When dsRNA has at least one unpaired nucleotide at the CC end, stability (and thus effective concentration in the cell) is improved interferon. The present sequence is that of a target DNA of the invention or content the concentration in the cells with the concentration in the cells with concentration in the cells with concentration.
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Best Local Sim
Matches. 25;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-683450/74.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-JAN-2001; 2001DE-01000588.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Inhibiting expression of target genes, useful e.g. for treating tumors, by introducing into cells two double-stranded RNAs that are complementary
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                                                                                                                                                                                                     Oligoribonucleotide; interferon; oncogene; cytokine;
                                                                                                                                                                                                                                                       Human
                                                                                                                                                                                                                                                                                                          23-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                           ABX09965 standard; DNA; 3840
                                                                                                                                                                           prion;
                                                                                                                                                                                                                                                                                                                                                            ABX09965;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1637
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                                                                                                                                                                                                                                                       mdr-1
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                                                                                                                                                                           inhibition;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CGAACCAGGGCACGTGCAATGGCGA 1613
                                                                                                                                                                                                                                                       DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 25; DB 6 ilarity 100.0%; Pred. No. 0.14; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                       fragment SEQ ID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                738 C;
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                                                                                                                                                                                                           Id;
                                                                                                                                                                                                           developmental;
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RESULT 11
ABL91687/c
ID ABL91687 standard; DNA; 3840
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              structure of not more than 49 consecutive nucleotides (nt), where at least a segment of one strand of the ds structure is complementary with the target gene and the cells are treated with interferon before introduction of dsRNAI. The method is used to inhibit expression of target genes, particularly oncogenes, cytokine genes, Id (not defined) protein genes; developmental or prion genes, or genes expressed in pathogenic organisms (particularly plasmodia) or in viruses or virolds (pathogenic in humans, animals or plants). Treating the cells with interferon greatly increases the extent to which dsRNA can inhibit expression of the target genes, and the effect is even greater when dsRNA are modified to increase their stability. ABX09936-ABX10075 represent gene fragments used to illustrate the method of the invention
               WPI; 2002-270454/32
                                          Kreutzer R,
                                                                                                    09-JAN-2001; 2001DE-01000586.
                                                                                                                                  09-JAN-2001; 2001DE-01000586
                                                                                                                                                                 11-APR-2002.
                                                                                                                                                                                             DE10100586-C1.
                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                       cytostatic; virucide; protozoacide; antibacterial; ds
                                                                                                                                                                                                                                                                    Human; HIV; HCV; gene expression; oligoribonucleotide; tumour; pathogen; Plasmodium; virus; viroid; cytokine; prion; antisense oligonucleotide;
                                                                                                                                                                                                                                                                                                                 Human polynucleotide SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                28-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                             ABL91687;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention describes a novel method for inhibiting expression of a target gene by introducing into the cell that contains the target gene at least one oligoribonucleotide (dsRNAI) that has a double-stranded (ds)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Inhibiting expression of target genes, e.g. oncogenes, in cells, by introduction of complementary double-stranded oligoribonucleotide, treating the cell with interferon.
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                                                                      (RIBO-)
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                                                                        RIBOPHARMA AG.
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ilarity 100.0%;
Conservative (
                                          Limmer S,
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                                        Hadwiger P;
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RESULT 12
ABQ78185/c
ID ABQ78185 standard; cDNA; 3843
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Best Local (
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                                                                                                                                                                  31-JAN-2001; 2001JP-00024723
                                                                                                                                                                                                                                                                                                                                                                                                                       Human;
single
                       Disclosure; Page 6-11; 14pp; Japanese
                                                    primer
                                                                Presuming the side effect
                                                                                             P-PSDB; ABB83950
                                                                                                          WPI; 2002-639348/69
                                                                                                                                                                                              31-JAN-2001; 2001JP-00024723.
                                                                                                                                                                                                                                                    JP2002223769-A.
                                                                                                                                                                                                                                                                                                          variation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABQ78185,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 3840 BP; 1130 A; 738 C; 957 G;
                                                                                                                                                                                                                          13-AUG-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 13; Page 28-30; 104pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         terminal bases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Inhibiting gene expression in cells, useful for e.g. treating tumors, introducing double-stranded complementary oligoRNA having unpaired
                                                                                                                                      (SRLS-) SRL
                                                      set.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25;
                                                                                                                                                                                                                                                                                                                                                                                                                     immunosuppressant; tacrolimus; cyclosporin; MDR1; SNP;
nucleotide polymorphism; gene; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CGAACCAGGGCACGTGCAATGGCGA 1613
                                                                                                                                                                                                                                                                                                                                                                                                                                                              encoding cDNA SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                            replace(2677,A/T)
/*tag= b
                                                                                                                                                                                                                                                                                                                       /product= "MDR1"
                                                                                                                                                                                                                                                                                                                                   /*tag=
                                                                                                                                                                                                                                                                             standard_name= "Single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                                location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                     .3843
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                                                                 of an
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 25;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ВÞ
                                                                 immunosuppressant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1015 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 3840;
                                                                 comprises
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The invention relates to presumption of the side effect of at least one

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ABT14496/c
ID ABT144
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AC ABT144
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PT 19-MAI
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PF 19-MAI
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PN WPI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 13
밁
                                                ફ
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Best Local S
Matches 25
                                                                                                          Query Match
Best Local S
Matches 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Macaca mulatta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rhesus monkey; s
P-glycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rhesus monkey P-glycoprotein coding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABT14496
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                                                                                                                                                                                                                                                                       The invention comprises the amino acid and coding sequence of a rhesus monkey (Macaca mulatta) P-glycoprotein and related P-glycoproteins. The DNA and protein sequences of the invention are useful in assays for evaluating the bioavailability of drugs, as well as the optimisation or discovery of drugs for the treatment of disease associated with P-glycoprotein transporter activity. The present cDNA sequence encodes the Rhesus monkey P-glycoprotein of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Isolated nucleic acid molecule encoding a P-glycoprotein of monkey, useful in assays for evaluating bioavailability of as for the optimization or discovery of drugs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-MAR-2001; 2001US-0277095P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-MAR-2002; 2002WO-US008325
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                                                                                                                                                                                                              Sequence 3852 BP; 1123 A; 750 C; 967 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3; Page 51-56; 103pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENTEST CORP
                                                                                                                                   Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard; cDNA; 3852
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CGAACCAGGGCACGTGCAATGGCGA 1616
                                                         CGAACCAGGGCACGTGCAATGGCGA 25
  CGAACCAGGGCACGTGCAATGGCGA 1625
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                                                                                                          100.0%;
llarity 100.0%;
Conservative 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gene; ss; P-glycoprotein inhibitor; drug bioavailability;
; P-glycoprotein transporter-related disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SR;
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Pred. No. 0.14;
                                                                                                                                         Score 25;
Pred. No.
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                                                                                                               Mismatches
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                                                                                                                                                                                                                          1012 T;
                                                                                                                                                                      BB
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                                                                                                                                                                Length 3852;
                                                                                                                                                                                                                            0 U; 0 Other;
                                                                                                                  Indels
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Best Matches Query Match

Local

Similarity

100.0%; Score 25; 100.0%; Pred. No.

0.14; DB 12; T; 0

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Gaps

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Conservative

Sequence 3858

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1132 A; 774

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984

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Other;

Length 3858; Indels

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ADH22599/c

ID ADH2259/x

AC ADH225

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YX I1-MAR

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XX INCOME

I Treat:

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                                                                               cc This invention relates to novel isolated polynucleotides identified as common transporters and ion channels (TRICH), and the encoded polypeptides thereof. Specifically, it describes using these TRICH molecules, as well cas agonists, antagonists, antibodies, expression vectors and host cells, in appropriate screening and toxicity assays to assess the effects of carterioscompounds on TRICH expression. The present invention describes TRICH compositions that are useful in the diagnosis, treatment and carterioscompounds of the carterioscompounds of the carterion as cell proliferative (e.g. arteriosclerosis, cancer), autoimmune/inflammatory (e.g. ALDS, asthma), carterioscompounds of the carterions accordingly, these TRICH compositions and for infections. Accordingly, these TRICH conjectures such as cytostatic, antinflammatory, immunosuppressive, cativities such as cytostatic, antinflammatory, immunosuppressive, contributed to a microarray is useful in monitoring or measuring proteingroties. This polymucleotide sequence is a human TRICH cDNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-MAY-2002;
03-MAY-2002;
05-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Baughn MR,
Emerling B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cDNA encoding a human transporter & ion channel (TRICH) protein SeqID97.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADH22599
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New human transporters and ion channels (TRICH), useful for diagnosing, treating and preventing diseases or conditions associated with the aberrant TRICH expression e.g. cancer, AIDS, atherosclerosis, epilepsy,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     neurological; epilepsy; stroke; developmental; Cushing's syndrome; hypothyroidism; infection; gene therapy; cytostatic; antiinflammatory; immunosuppressive; antiasthmatic; anticonvulsant; nootropic; neuroprotective.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11-MAR-2004
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P-PSDB; ADH22533.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ng BM, Griffin JA, F
Kable AE, Khare R,
ar J, Richardson TW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard; cDNA;
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; 2002US-0386497P.
; 2002US-0388180P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97; 448pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hafalia AJA, Ison CH
, Lee SY, Lee S, Ma
, Swarnakar A, Tran
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AJA, Ison CH, Jackson
, Lee S, Mason PM, Ma
kar A, Tran UK, Chawla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            channel;
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Chawla NK;
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RESULT 15
AAZ49332/c
ID AAZ49332 standard; cDNA; 3860 BP.
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CC This sequence represents cDNA encoding human wild-type multidrug CC responsible for the export of drugs from cells, particularly cancer cells. Wild-type MDR-1 shows increased resistance to etoposide and CC decreased resistance to vinca alkaloide. compared with a mutant form CC (AAYS6187) where the Gly at position 185 is replaced by Val. The CC invention relates to transducing haematopoietic stem cells with nucleic conditied haematopoietic stem cells are useful in bone marrow conditied haematopoietic stem cells are useful in bone marrow transplantation (to reconstitute haematopoietic systems in patients who can be undergone chemotherapy or radiation therapy) and in ex vivo gene convertible, e.g., thalassaemia, Gauchier's disease, sickle cell anaemia or cells, e.g., thalassaemia, Gauchier's disease, sickle cell anaemia or cells, e.g., thalassaemia, Gauchier's disease, sickle cell anaemia or cells, e.g., thalassaemia, Gauchier's disease, sickle cell anaemia or cells, e.g., thalassaemia, Gauchier's disease, sickle cell anaemia or cells, e.g., thalassaemia, Gauchier's disease, sickle cell anaemia or cells, e.g., thalassaemia, Gauchier's disease, sickle cell anaemia or cells, e.g., thalassaemia, Gauchier's disease, sickle cell anaemia or cells, e.g., thalassaemia, Gauchier's disease, sickle cell anaemia or cells, e.g., thalassaemia, Gauchier's disease, sickle cell anaemia or cells, e.g., thalassaemia, Gauchier's disease, sickle cell anaemia or cells, e.g., thalassaemia, Gauchier's disease, sickle cell anaemia or cells, e.g., thalassaemia, Gauchier's disease, sickle cell anaemia or cells that express MDR-1 will be protected against chemotherapy. Expansion of (rare) comparated while the patient is undergoing chemotherapy. Expansion of (rare) comparated biochemical analysis. Overexpression of MDR-1 allows cytokine-driven
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Multidrug resistance; MDR-1; P-glycoprotein; transmembrane efflux pump; haematopoietic stem cell; transduction; bone marrow transplantation; chemotherapy; radiation therapy; cancer; gene therapy; gene replacement genetic defect; thalassaemia; Gauchier's disease; sickle cell anaemia; leukaemia; ex vivo expansion; cytokine; wild-type; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 10; Page 68-70; 113pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ex vivo expansion of hematopoietic stem cells transduced with a sequence encoding human multidrug resistance-1, used for bone marrow
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sorrentino B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28-MAY-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human wild-type multidrug resistance-1 (MDR-1) cDNA.
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/note= "cDNA sequence of G185V human mutant MDR-1 given
in AAZ49333"
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                                            Matches
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Best Local
                                                                                       Sequence 3860 BP; 1135 A; 746 C; 957 G;
                                                                                                            expansion of haematopoietic stem cells by at least 10-fold compared with a maximum of 4-fold in known procedures
1640
                                            25;
                     1 CGAACCAGGGCACGTGCAATGGCGA 25
                                                       Similarity
CGAACCAGGGCACGTGCAATGGCGA 1616
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Pred. No.
                                                                                         1022
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                                                                  Length 3860;
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Search completed: February 9, 2005, 16:05:59 Job time : 246.69 secs

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Run on:
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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1: /cgn2_6/ptodata/1/pubpna/US07
2: /cgn2_6/ptodata/1/pubpna/PCT
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Copyright (c) 1993 - 2005 Compugen Ltd.
/cgn2 6/ptodata/1/pubpna/USO7_PUBCOMB.seq:*
/cgn2 6/ptodata/1/pubpna/USO7_NEW_PUB.seq:*
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/cgn2 6/ptodata/1/pubpna/USO9_PUBCOMB.seq:*
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(without alignments)
539.601 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIE

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C 5 25 100.0 3860
C 6 25 100.0 3924
C 8 25 100.0 3924
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RESULT 2 US-10-363-112-48/c ; Sequence 48, Application US/10363112

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ω | 32 | 31 | 30 | 29 | 28 | 27 | 26 | 25 | 24 | 23 | 22 | 21 | 20 | 19 | 18 | 17 | 16 | 15 | 14 | 13 | 12 |
| 18.6 | 18.6 | 18.6 | 18.6 | 18.8 | 19 | 19.2 | 19.2 | 19.2 | 19.2 | 19.2 | 19.2 | 20.2 | 20.2 | 20.2 | 20.2 | 20.2 | 20.2 | 20.2 | 20.2 | 20.2 | 21.8 | 21.8 | | | | | | 25 | ა
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| 74.4 | 74.4 | 74.4 | 74.4 | 75.2 | | 80 | 8 | В | 80 | 76.8 | 76.8 | 80.8 | 80.8 | 80.8 | 80.8 | 80.8 | 80.8 | 80.8 | 80.8 | 80.8 | 87.2 | 87.2 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | • | 100.0 | 100.0 | • | 100.0 | 100.0 |
| 2021 | 1940 | 1818 | 1175 | 483 | 19 | 2731748 | 55001 | 55001 | 7615 | 747 | 711 | 4788 | 4425 | 4369 | 4298 | 4254 | 4254 | 4254 | 4189 | 3912 | 4317 | 4317 | 8630 | 8630 | 4669 | 4646 | 4646 | 4646 | 4646 | 4646 | 4643 | 4643 | 4643 |
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| US-09-873-409-15 | US-09-873-409-16 | US-10-156-761-2443 | _ | US-10-369-493-29889 | US-10-016-490C-38 | S | US-10-348-750-4 | US-10-160-497-4 | US-10-004-113-59 | US-10-369-493-41419 | 10-369-493-3318 | US-09-866-866A-7 | US-09-769-097-3 | US-09-769-097-1 | US-10-335-053-32 | US-10-152-319A-1484 | US-10-388-934-265 | US-09-917-800A-1424 | US-09-866-866A-5 | US-09-917-800A-1560 | | US-10-044-671-1 | US-09-306-417-2 | US-09-306-417-1 | US-10-680-516-1 | US-10-775-169-198 | , - | US-10-641-643-1167 | US-09-968-007A-747 | US-09-968-007A-459 | US-10-007-926A-258 | | 621- |
| Sequence 15, Appr | • | 6 24 | Sequence II, Appr | Sequence 29889, A | Sequence 38, Appr | , ·· | ••• | 4. | Sequence 59, Appi | | SSIEG, | /, Appii | ٠, | | e 32, | Sequence 1484, Ap | 265 | 424 | Sequence 5, Appli | Sequence 1560, Ap | Sequence 1, Appli | , ,- | Sequence 2, Appli | | Sequence 1, Appii | | | ۳. | 747, | equence 459, | | 1, At | Sequence 2, Appli |

ALIGNMENTS

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; LOCATION: (1)..(3825)
US-10-363-112-48
                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: PCT/EP02/00152
PRIOR PILING DATE: 2002-01-09
PRIOR PPLICATION NUMBER: DE 1010586.5
PRIOR PILING DATE: 2001-01-09
PRIOR EILING DATE: 2001-10-09
PRIOR APPLICATION NUMBER: DE 10155280.7
PRIOR PILING DATE: 2001-10-26
PRIOR APPLICATION NUMBER: DE 10158411.3
PRIOR PILING DATE: 2001-11-29
PRIOR PILING DATE: 2001-11-07
NUMBER: OF SEQ ID NOS: 173
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                                                                                Query Match
Best Local S
Matches 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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Best Local Similarity
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                                                                                                                                                                   -10-384-339C-30
                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Homo sapiens
PUBLICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Kreutzer, Roland
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR INHIBITING EXPRESSION OF A TARGET GE
FILE REFERENCE: 20200/2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: PCT/AU01/01093
PRIOR FILING DATE: 2001-08-31
PRIOR APPLICATION NUMBER: 60/229,663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: 007643-0302189
CURRENT APPLICATION NUMBER: US/10/363,112
CURRENT FILING DATE: 2003-11-03
                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/10/384,339C CURRENT FILING DATE: 2003-03-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE: 2000-08-31
NUMBER OF SEQ ID NOS: 62
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APPLICANT: HARRIS, MATTHEW
TITLE OF INVENTION: MODIFIED PROTEINS, ISOLATED NOVEL PEPTIDES, AND USES THEREOF
                                                                                                                                                                                     TITLE: mdr-1
PATENT DOCUMENT NUMBER: AF016535
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
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1637 CGAACCAGGGCACGTGCAATGGCGA 1613
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Conservative 0
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100.0%; Pred. No. 0.035;
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Pred. No. 0.035;
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; TYPE: DNA
; ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 1, Application US/09866866A Patent No. US20020102244A1
GENERAL INFORMATION:
                                                                                  Matches
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                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                              APPLICANT: Sorrentino, Brian
APPLICANT: Schuetz, John
TITLE OF INVENTION: A Method of Identifying and/or Isolating Stem Cells
FILE REFERENCE: 1340-1-021CIP2
CURRENT APPLICATION NUMBER: US/09/866,866A
CURRENT FILING DATE: 2001-08-30
PRIOR APPLICATION NUMBER: 09/584,586
PRIOR APPLICATION NUMBER: 09/584,586
PRIOR APPLICATION NUMBER: PCT/US99/11825
PRIOR APPLICATION NUMBER: PCT/US99/11825
PRIOR FILING DATE: 1999-05-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Hanscom, Sara
APPLICANT: Cresp1, Charles
TITLE OF INVENTION: P-GLYCOPROTEINS AND USES THEREOF
FILE REFERENCE: G00307/70019
FILE REFERENCE: G00307/70019
CURRENT FAPPLICATION NUMBER: US/10/101,433A
CURRENT FILLING DATE: 2002-03-19
PRIOR APPLICATION NUMBER: US 60/277,095
                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/086,988 PRIOR FILING DATE: 1998-05-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 2001-03-19 NUMBER OF SEQ ID NOS: 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Macaca mulatta
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: CDS
LOCATION: (1)..(3852)
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                                                                                                      Local
1640 CGAACCAGGGCACGTGCAATGGCGA 1616
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                                                                                  25;
                                                                                                      Similarity
                                CGAACCAGGGCACGTGCAATGGCGA 25
                                                                                100.0%; Score 25; DB 9; llarity 100.0%; Pred. No. 0.035; Conservative 0; Mismatches
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100.0%; Pred. No.
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                                                                                                                       DB 9;
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                                                                                                                         Length 3860;
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US-09-866-866A-3/c
, Sequence 3, Application US/09866866A
; Patent No. US20020102244A1
; GENERAL INFORMATION:

APPLICANT: Sorrentino, Brian

RESULT 6

GENE EXPRESSION

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APPLICANT: HOTTE, Darci T.

APPLICANT: Vockley, Joseph G.

APPLICANT: Scherf, Uwe
APPLICANT: Scherf, Uwe
FILE Scherf, Uwe
FILE OF INVENTION: Gene Expression Profiles in Liver Cancer
FILE REFERENCE: 44921-5028-W0
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT FILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
NUMBER OF SEQ ID NOS: 3950
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CURRENT APPLICATION NUMBER: US/09/866,866A
CURRENT FILING DATE: 2001-08-30
PRIOR APPLICATION NUMBER: 09/584,586
PRIOR FILING DATE: 2000-05-31
PRIOR APPLICATION NUMBER: PCT/US99/11825
PRIOR APPLICATION NUMBER: PCT/US99/11825
PRIOR FILING DATE: 1998-05-27
PRIOR FILING DATE: 1998-05-28
NUMBER OF SEQ ID NOS: 27
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US-10-641-643-1168/c
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                                                                                  Sequence 1168, Application US/10641643 Publication No. US20040077003A1 GENERAL INFORMATION:
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SEQ ID NO 2299
LENGTH: 3924
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA ORGANISM: homo mapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Genbank Accession No. US20020142981A1 M23234
                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
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                                                                                                                                                                                                                                       1678
Susan G. Stuart
Jeffrey J. Seilhamer
TITLE OF INVENTION: COMPOSITION FOR THE DETECTION OF BLOOD CELL
                                                                  APPLICANT: Cocks, Benjamin G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 CGAACCAGGGCACGTGCAATGGCGA 25
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                                                                                                                                                                                                                                                                                                                     100.0%; Score 25; DB 9; Length 3924; 100.0%; Pred. No. 0.035; tive 0; Mismatches 0; Indels (
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US-10-619-359A-1/c
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                                                                                                                                                   FILE REFERENCE: G00307.70020.US
CURRENT APPLICATION NUMBER: US/10/619,359A
CURRENT FILING DATE: 2003-07-14
PRIOR APPLICATION NUMBER: US 60/156,921
PRIOR FILING DATE: 1999-09-28
PRIOR APPLICATION NUMBER: US 60/158,818
PRIOR APPLICATION NUMBER: US 09/672,810
PRIOR APPLICATION NUMBER: US 09/672,810
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: STOCKER, PENNY J.
APPLICANT: STEIMEL-CRESPI, DOROTHY T.
APPLICANT: CRESPI, CHARLES L.
TITLE OF INVENTION: P-GLYCOPROTEINS AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 1168:
SEQUENCE CHARACTERISTICS:
LENGTH: 3924 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                       SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                NUMBER OF SEQ ID NOS:
ORGANISM: Macaca fascicularis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Word Perfect 6.1 for Windows/MS-DOS 6.2
SOPTWARE: Word Perfect 6.1 for Windows/MS-DOS 6.2
APPLICATION UNMBER: US/10/641,643
EILING DATE: 14-Aug-2003
CLASSIFICATION UNMBER: -CUNKNOWN>
PRIOR APPLICATION OATA:
APPLICATION UNBER: -CUNKNOWN>
FILING DATE: -CUNKNOWN>
ATTORNEY/AGENT INFORMATION:
NAME: Zeller, Karen J.
REGISTRATION UNMBER: 37,071
REFERENCE/DOCKET NUMBER: PA-0001 US
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                    ENGTH:
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CORRESPONDENCE ADDRESS:
CORRESPONDENCE INCYTE PHARMACEUTICALS, INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLONE: 9187501
SEQUENCE DESCRIPTION: SEQ ID NO: 1168:
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US-09-805-020-30/c

; Sequence 30, Application US/09805020

; Publication No. US20020086384A1

; GENERAL INFORMATION:

; APPLICANT: LEVINE, Zurit

; TITLE OF INVENTION: SPITICE VARIANTS OF ONCOGENES

; FILE REFERENCE: 2786-0168P

; CURRENT APPLICATION NUMBER: US/09/805,020

; CURRENT FILING DATE: 2001-03-13

; NUMBER OF SEQ ID NOS: 72

; SOFTWARE: Patentin Ver. 2.1

; SOFTWARE: Patentin Ver. 2.1
US-09-805-020-30
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Best Local S
Matches 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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Best Local :
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APPLICANT: STOCKER, PENNY J.
APPLICANT: STEMMEL-CRESPI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: STEIMEL-CRESPI, DOROTHY T.
APPLICANT: CRESPI, CHARLES L.
TITLE OF INVENTION: P-GLYCOPPOTEINS AND USES THEREOF
FILE REFERENCE: G00307.70020.US
CURRENT APPLICATION NUMBER: US/10/619,359A
CURRENT FILING DATE: 2003-07-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 60/156,921
PRIOR FILING DATE: 1999-09-28
PRIOR RPLICATION NUMBER: US 60/158,818
PRIOR FILING DATE: 1999-10-12
PRIOR APPLICATION NUMBER: US 09/672,810
PRIOR PILING DATE: 2000-09-28
NUMBER OF SEQ ID NOS: 18
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                 NAME/KEY: misc_feature
LOCATION: (1)...(4533)
OTHER INFORMATION: any n =
                                                                      TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Macaca fascicularis
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LOCATION: (100)...(3949)
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                                                                                                                              ENGTH: 4533
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25; Conserv
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25; Conservative
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(100)...(3940)
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100.0%; Pred. No. 0.035;
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               a,c,g,t any unknown or other
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Pred. No. 0.035;
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; ORGANISM: Homo sapiens
US-10-072-621-2
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US-10-072-621-2/c
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Publication No. US20020169137A1
GENERAL INFORMATION:
APPLICANT: Reiner, Peter B.
APPLICANT: Connop, Bruce P.
APPLICANT: Pollard, Michelle
TITLE OF INVENTION: REGULATION OF AMYLOID PRECURSOR PROTEIN EXPRESSION
TITLE OF INVENTION: BY MODIFICATION OF ABC TRANSPORTER EXPRESSION OR ACTIVITY
FILE REFERENCE: 100103,402
CURRENT APPLICATION NUMBER: US/10/072,621
CURRENT FILING DATE: 2002-02-08
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            TITLE OF INVENTION: Nucleic Acid Molecules and Proteins For The Identification, TITLE OF INVENTION: Assessment, Prevention, and Therapy of Ovarian Cancer FILE REFERENCE: MRI-030 CURRENT APPLICATION NUMBER: US/10/097,340 CURRENT FILING DATE: 2002-03-14 PRIOR APPLICATION NUMBER: 60/276,025 PRIOR FILING DATE: 2001-03-14 PRIOR APPLICATION NUMBER: 60/25,149 PRIOR APPLICATION NUMBER: 60/325,149 PRIOR FILING DATE: 2001-09-26 PRIOR FILING DATE: 2001-09-26 PRIOR FILING DATE: 2001-09-26 PRIOR FILING DATE: 2001-03-14 PRIOR APPLICATION NUMBER: 60/276,026 PRIOR FILING DATE: 2001-03-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: John MONAHAN
APPLICANT: Manjula GANI
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APPLICATION NUMBER: 60/324,967
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Sebastian HOERSCH
Shubhangi KAMATKAR
Steve G. KOVATS
Rachel E. MEYERS
Michael MORRISEY
Peter OLANDT
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Karen GLATT
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Robert C. BAST, Jr.
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RESULT 15
US-09-968-007A-459/c
US-09-968-007A-459/c
; Sequence 459, Application US/09968007A
; Publication No. US20040115625A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     , OTHER INFORMATION: atp-binding cassette, sub-family b , OTHER INFORMATION: (mdr/tap), member 1 (ABCB1) gene. US-10-007-926A-258
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US-10-007-926A-258/c
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-097-340-1
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PRIOR APPLICATION NUMBER: 60/311,732
PRIOR FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: 60/325,102
PRIOR FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 60/323,580
PRIOR FILING DATE: 2001-09-19
PRIOR FILING DATE: 2001-09-19
PRIOR FILING DATE: 2001-09-19
PRIOR FILING DATE: 2001-09-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 258, Application US/10007926A Publication No. US20030143539A1 GENERAL INFORMATION:
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Best Local Similarity 100.0%; Pred. No. 0.035;
Matches 25; Conservative 0; Mismatches
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Best Local Similarity
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APPLICANT: NGUYEN, CATHERINE
APPLICANT: VIENS, PATRICE
APPLICANT: VIENS, PATRICE
APPLICANT: VIENS, PATRICE
APPLICANT: FERT, VINCENT
TITLE OF INVENTION: GENE EXPRESSION PROFILING OF PRIMARY BREAST CARCINOMAS
TITLE OF INVENTION: USING ARRAYS OF CANDIDATE GENES
FILE REFERENCE: 1546-R-00
CURRENT APPLICATION NUMBER: US/10/007,926A
CURRENT FILING DATE: 2001-12-07
PRIOR APPLICATION NUMBER: 60/254,090
PRIOR FILING DATE: 2001-12-08
NUMBER OF SEQ ID NOS: 468
SOFTWARE: Patentin Ver. 2.1
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APPLICANT: HOULGATTE, REMI
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APPLICANT: Ebner, Reinhard TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Usining Signa TITLE OF INVENTION: Gene Sets FILE REFERENCE: 689290-11 CURRENT APPLICATION NUMBER: US/09/968,007A CURRENT FILING DATE: 2001-10-02
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BIRNBAUM, DANIEL
NGUYEN, CATHERINE
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PRIOR APPLICATION NUMBER: US/60/237,294
PRIOR APPLICATION NUMBER: US/60/237,295
PRIOR FILING DATE: 2000-10-02
PRIOR PRIOR PRICATION NUMBER: US/60/237,316
PRIOR APPLICATION NUMBER: US/60/237,316
PRIOR PILING DATE: 2000-10-02
PRIOR PILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 1001
SOFTWARE: PATENTIN VETBION 3.0
SEQ ID NO 459
IENGTH. 4646
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-968-007A-459
Search completed: February Job time : 268.667 secs
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Local Similarity 100.0%; Pred. No. 0.035;
hes 25; Conservative 0; Mismatches
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CC475658 CC240_301
AK050345 Mus muscu
AK030328 Mus muscu
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BP220580
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AV191114 AV191114
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AY408955 Pan trogl
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AG179442 Pan trogl
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F08704 HSCZOA021 n
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| | 44 | 4 | 42 | 41 | 40 | 39 | 38 | 37 | 36 | 35 | 34 | 33 | 32 | 31 | 30 | 29 | 28 | 27 | 26 | 25 |
| 18.6 | 18.6 | 18.8 | 18.8 | 18.8 | 18.8 | 18.8 | 18.8 | 18.8 | 18.8 | 18.8 | 18.8 | 19.2 | 19.2 | 19.2 | 19.2 | 19.2 | 19.2 | 19.2 | 19.2 | 19.2 |
| 74.4 | 74.4 | 75.2 | 75.2 | 75.2 | 75.2 | 75.2 | 75.2 | 75.2 | 75.2 | 75.2 | 75.2 | 76.8 | 76.8 | 76.8 | 76.8 | 76.8 | 76.8 | 76.8 | 76.8 | 76.8 |
| 375 | 360 | 859 | 800 | 757 | 710 | 656 | 641 | 639 | 638 | 580 | 392 | 3721 | 845 | 754 | 726 | 663 | 578 | 569 | 446 | 429 |
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| C63703 | C67438 | BG704400 | CN816473 | BU314758 | AA185993 | BJ818135 | BJ814960 | CF599341 | AA175587 | BE687194 | AI115290 | BC032414 | BG818192 | CN431066 | BH112888 | CN431062 | CE805931 | CR241012 | BQ343554 | CF136217 |
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ALIGNMENTS

JOURNAL MEDLINE PUBMED COMMENT REFERENCE AUTHORS SOURCE ORGANISM VERSION KEYWORDS RESULT 1 F07318 LOCUS ACCESSION FEATURES DEFINITION TITLE source Homo sapiens Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 290) 1 (bases 1 to 290) 1 (bases 1 to 290) 2 (bases 1 to 290) 1 (bases 1 to 290) 2 (bases 1 to 290) 2 (bases 1 to 290) 3 (bases 1 to 290) 4 (bases 1 to 290) 2 (bases 1 to 290) 3 (bases 1 to 290) 4 (bases 1 to 290) 5 (bases 1 to 290) 4 (bases 1 to 290) 5 (bases 1 to 290) 6 (bases 2 to 290) 6 (bases 2 to 290) 6 (bases 1 to 290) 7 (bases 1 to 290) 7 (bases 1 to 290) 8 (bases 1 to 290) 1 (bases 1 to 290) Genexpress library idt: C; Genexpress sequence_idt: y1c-22a07 Seq primer: (-21)MI3_universal. Location/Qualifiers Genethon Centre de recherch 1, rue de l'Internationale, Tel: 33169472800 Fax: 33160778698 F07318 290 bp mRNA linear HSC22A071 normalized infant brain cDNA Homo sapiens and F07318.1 c-22a07, mRNA sequence. Single read Genexpress-Genethon Contact: Genethon 95277534 C. R. Acad. Sci. III, Homo sapiens (human) 7757816 its expression genexpress@genethon.fr /tissue_type="total brain" /dev_gtage="3 months old" /clome_lib="normalized infant brain cDNA" /note="Organ: brain; Vector: lafmid BA; Site_1: HindIII; /site_2: NotI; sex=Female; dev_stage=3 months old; isolate=muscular atrophy patient; tissue_type=total brain; total mRNA was oligo-(dT) primed and directionally cloned 5' -> 3' into the HindIII -> NotI sites of the /organism="Homo sapiens" /mol type="mRNA" /db_xref="taxon:9606" /clone="c-22a07" GI:672970 'sex="Female" recherche Sci. Vie 318 (2), 263-272 (1995) 290 bp ne sur le Ge BP60 91002 mRNA Genome Humain 02 EVRY Cedex, linear FRANCE cDNA clone EST 20-FEB-1995

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25; Conserv
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Genethon Centre de recherche sur le Genome Humain
1, rue de l'Internationale, BP60 91002 EVRY Cedex,
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1 (bases 1 to 315)

Auffray,C., Behar,G., Bois,F., Bouchier,C., da Silva,C.,

Aufgray,M.D., Duprat,S., Houlgatte,R., Jumeau,M.N., Lamy,B.,

Devignes,M.D., Duprat,S., Houlgatte,R., Jumeau,M.N., Lamy,B.,

Lorenzo,F., Mitchell,H., Mariage-Samson,R., Pietu,G., Pouliot

Sebastiani-Kabaktchis,C. and Tessier,A.

IMAGE: molecular_integration of the analysis of the human gen
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Seq_primer: (-21)MI3_universal.
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                                      CGAACCAGGGCACGTGCAATGGCGA 25
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llarity 100.0%;
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/dev_stage="3 months old"
/clome_libb="normalized infant brain cDNA"
/clome_libb="normalized infant brain cDNA"
/note="Organ: brain; Vector: lafmid BA; Site_1: HindIII;
Site_2: NotI; sex=Female; dev_stage=3 months old;
isolate=muscular atrophy patient; tissue_type=total
brain; total mRNA was oligo-(dT) primed and directionally
cloned 5' -> 3' into the HindIII -> NotI sites of the
lafmid BA vector. Clone library from B.Soares, Psychiatry
Dept. Columbia University, USA. Normalization_method:
Bento Soares, P.N.A.S in press"
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mol_type="mRNA"
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HSC11E051 normalized infant
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Genexpress Genethon centre de recherche sur le Genome Humain
Genethon Centre de recherche sur le Genome Humain
Genethon Centre de PRO 91002 EVRY Cedex,
Homo sapiens (human)
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Seq primer: (-21)M13_universal.
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Auffray,C., Behar,G., Bois,F., Bouchier,C., da Silva,C.,
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/dev stage="3 months old"
/clone_lib="normalized infant brain cDNA"
/clone_lib="normalized infant brain dBA; Site_1: HindIII;
/note="Organ: brain; Vector: lafmid BA; Site_1: HindIII;
Site_2: NOLI; sex=Female; dev stage=3 months old;
isolate=muscular atrophy patient; tissue_type=total
brain; total mRNA was oligo-(dT) primed and directionally
cloned 5' -> 3' into the HindIII -> NotI sites of the
lafmid BA vector. Clone library from B.Soares, Psychiatry
Dept. Columbia University, USA. Normalization_method:
Bento Soares, P.N.A.S in press"
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/db_xref="taxon:9606"
/clone="c-zoc01"
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Best Local Similarity
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1 (bases 1 to 328)
2 (bases 1 to 328)
2 (bases 1 to 328)
3 (bases 1 to 328)
3 (bases 1 to 328)
4 (bases 1 to 328)
5 (bases 1 to 328)
5 (bases 1 to 328)
6 (bases 1 to 328)
7 (bases 1 to
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Seq_primer: (-21)MI3_universal.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Genexpress-Genethon
Genethon Centre de recherche sur le Genome Humain
1, rue de l'Internationale, BP60 91002 EVRY Cedex,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Fax: 33160778698
Human Genome Center, L-452
Lawrence Livermore National Laboratory
Livermore CA 94550
                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 357) Ghiso,N.S., Eveleth,G.G., Lieuallen,K. and Lennon,G.G. Infant brain cDNAs
                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens (human)
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T26505.1 GI:773822
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AB282E2R Infant brain, LLNL array of Dr. M. Soares 1NIB Homo
Bapiens cDNA clone LLAB282E2 5', mRNA sequence.
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                                                                                                     Contact: Greg G. Lennon
                                                                                                                                           7490096
                                                                                                                                                                                                                      Genomics 28 (3), 570-572
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/note="Organ: brain; Vector: lafinid BA; Site_1: HindIII;
Site_2: NotI; sex=Female; dev_stage=3 months old;
isolate=muscular atrophy patient; tissue type=total
brain; total mRNA was oligo-(dT) primed and directionally
cloned 5' -> 3' into the HindIII -> NotI sites of the
lafinid BA vector. Clone library from B. Soares, Psychiatry
Dept. Columbia University, USA. Normalization_method:
Bento Soares, P.N.A.S in press"
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/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="c-11e05"
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dev_stage="3 months old"
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Pred. No. 0.95;
Mismatches
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Best Local :
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Fax: 510 422 2282
Email: info@image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 469)

Hillier, L., Clark, N., Dubuque, T.; Elliston, K., Hawkins, M., Holman, M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Marra, M., Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trevakis, E., Waterston, R., Williamson, A., Wohldmann, P. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R53330 A69 bp mRNA linear EST 18-MAY-19
yg83b02.r1 Soares infant brain 1NIB Homo sapiens cDNA clone
IMAGE:39920 5' similar to gb:M23234 MULTIDRUG RESISTANCE PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25;
                                                                                                                                                                                                                                                                   Email: estewatson.wustl.edu
Insert Size: 2097
High quality sequence stops: 371 Source: IMAGE Consortium, LLNL
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Insert Length: 2097 Std Error: 0.00
Seq primer: M13RP1
                                                                                                                                                                                                                                                                                                                                                                                                                   Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. 1 Tel: 314 286 1800 Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (1995)
Contact: Wilson RK
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                                                                                                                                                                                                                                                  quality sequence stop: 371.
                                                                                                    /mol type="mRNA"
/db_xref="GDB:412461"
/db_xref="taxon:9606"
/clone="IMAGE:39920"
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/db_xref="taxon:9606"
/clone="LLAB282E2"
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/mol_type="mRNA"
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/dev_stage="73 days post natal"
//ab host="pH10B (ampicillin resistant)"
/clone lib="Soares infant brain INIB"
/note="Organ: whole brain; Vector: Lafmid BA; Site_1: Not
                                                                                  /sex="female"
                                                                                                                                                                                       'organism="Homo sapiens'
                                                                                                                                                                                                                                ocation/Qualifiers
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                                                        BP220580 Sugano cDNA library, COL03695, mRNA sequence.
BP220580
Homo
                                     BP220580.1 GI:52093485
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                                                                                                                                                                                                                                    25;
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1 (bases 1 to 568)
Suzuki, Y., Yamashita, R., Shirota, M., Sakakibara, Y., Chiba, J., Mizushima-Sugano, J., Nakai, K. and Sugano, S.
Sequence comparison of human and mouse genes reveals a homologous block structure in the promoter regions
Genome Res. 14 (9), 1711-1718 (2004)
Contact: Yutaka Suzuki
Department of Virology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP243819 Sugano cDNA library,
HEP21483, mRNA sequence.
BP243819
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Institute of Medical Science, University of Tokyo 4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan Email: yeuzukideme u-tokyo.ac.jp.
Location/Qualifiers
                                                                                                                                                                                                                                                 Similarity
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             sapiens (human)
sapiens
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/cell_line="HepG2"
/clone_lib="Sugano_cDNA_library, liver_HepG2"
/note="hepatoma"
                                                                                                                                                                                                                                                                                                                                                              /mol_type="mRNA"
/db_xref="taxon:9606"
/clone="HEP21483"
                                                                                                                                                                                                                                                                                                                                                                                                          organism="Homo sapiens"
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Pred. No. 0.99;
                                                                                                                                                                                                                                   Mismatches
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                                                                                    , colon
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                                                                                   Homo sapiens cDNA clone
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AG179058/c
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AG179058.1
GSS.
                                                                                                                                                                                                                        Submitted (02-AUG-2001) Asao Fujiyama, The Institute of Physical and Chemical Research (RIKEN), Genomic Sciences Center (GSC); 1-7-22 Suehiro-chou, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan (B-mail:chimpbesegsc.riken.go.jp, URL:http://hgp.gsc.riken.go.jp/, Tel:81-45-503-9111, Fax:81-45-503-9170)
Clones are derived from the chimpanzee BAC library RPCI-43 This BAC end was generated during the R&D process and may have higher chance of clone tracking errors.
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1 (bases 1 to 570)

Suzuki,Y., Yamashita,R., Shirota,M., Sakakibara,Y., Chiba,J., Mizushima-Sugano,J., Nakai,K. and Sugano,S.

Sequence comparison of human and mouse genes reveals a homologous block structure in the promoter regions
Genome Res. 14 (9), 1711-1718 (2004)

Contact: Yutaka Suzuki
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Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims_u-tokyo.ac.jp.
                                                                                                                                                                                                                                                                                                                                                                                                                Fujiyama,A., Hattori,M., Toyoda,A., Taylor,T.D., Yada,T.,
Totoki,Y., Watanabe,H. and Sakaki,Y.
Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pan troglodytes (chimpanzee)
Pan troglodytes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AG179058 669 bp DNA linear Pan troglodytes DNA, clone: RP43-051D12.T7, genomic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished
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Mammalia; Eutheria; Primates; Catarrhini, Hominidae; Pan.
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                                                                                              Vector : pBACe3.6
R.Site 1 : EcoRI
R.Site 2 : EcoRI.
Location/Qualifiers
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/organism="Pan troglodytes"
/mol_type="genomic DNA"
/db_xref="taxon:9598"
/clone="RP43-051D12.T7"
                                                                                   1. .669
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/clone_Tib="Sugano cDNA library, colon"
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/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="COL03695"
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1. .570
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Pred. No. 1;
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RESULT 10
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                      RESULT 11
AY408954/c
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Best Local Similarity
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Homo sapi
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Clones are derived from the chimpanzee BAC library RPCI-43 This BAC end was generated during the R&D process and may have higher chance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fujiyama, A., Hattori, M., Toyoda, A., Taylor, T.D., Yada, T., Totoki, Y., Watanabe, H. and Sakaki, Y.
Direct Submission
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Totoki,Y., Watanabe,H. and Sakaki,Y.
BAC end sequences of Library RPCI-43
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R.Site 2
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sapiens ABCB1 gene,
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/cell_type="lymphocytes"
/clone_llb="RPCI-43 Chimpanzee Male BAC Library"
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/db_xref="taxon:9598"
/clone="RP43-051L12.T7"
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/clone_lib="RPCI-43 Chimpanzee Male BAC Library"
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 3843 bp DNA linear GSS 15-DEC VIRTUAL TRANSCRIPT, partial sequence,
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     Direct Submission
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1640 CGAACCAGGGCACGTGCAATGGCGA 1616
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1 (bases 1 to 3843)
Clark, A.G., Glanowski, S., Nielson, R., Thomas, P., Kejariwal, A., Todd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F., Murphy, B., Ferriera, S., Wang, G., Zheng, X.H., White, T.J., Sninsky, J.J., Adams, M.D. and Cargill, M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 CGAACCAGGGCACGTGCAATGGCGA 25
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Direct Submission
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Science 302 (5652), 1960-1963 (2003)
2 (bases 1 to 3843)
Clark, A.G., Glanowski, S., Nielson, R., Thomas, P., Kejariwal, Todd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F., Murphy, B., Ferriera, S., Wang, G., Zheng, X.H., White, T.J., Sninsky, J.J., Adams, M.D. and Cargill, M.
                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Pan.
1 (bases 1 to 384)
Clark, A.G., Glanowski, S., Nielson, R., Thomas, P., Kejariwal, A., Todd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F., Murphy, B., Ferriera, S., Wang, G., Zheng, X.H., White, T.J., Sninsky, J.J., Adams, M.D. and Cargill, M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pan troglodytes ABCB1 gene, VIRTUAL TRANSCRIPT, partial genomic survey sequence.
AY408955
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                         Pan troglodytes (chimpanzee)
Pan troglodytes
                                                                                                                                Science 302 (5652), 1960-1963 (2003) 14671302
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                                                                                                                                                                                                               Inferring nonneutral evolution from human-chimp-mouse orthologous
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequence was made by sequencing based on alignment.
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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/locus_tag="HCM3396"
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                                                                                    Kejariwal, A.,
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BC042531/c
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                                                                                                                                                                                           Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LINL at: http://image.llnl.gov Series: IRAK Plate: 72 Row: 1 Column: 3
This clone was selected for full length sequencing because it passed the following selection criteria: matched mRNA gi: 9961253
This clone has the following problem: frame shifted.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Direct Submission
Submitted (02-JAN-2003) National Institutes of Health, Mammalian Gene Collection (MGC), Cancer Genomics Office, National Cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              #CO42531 4035 bp π
Homo Bapiens, ATP-binding cassette,
4, clone IMAGE:4837724, mRNA.
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                                                                                                                                                                                                                                                                                                                                                                       Web site: http://www.hgsc.bcm.tmc.edu/cdna/
Contact: amg@bcm.tmc.edu
Gunaratne, P.H., Garcia, A.M., Lu, X., Hulyk, S.W., Loulseged,
Gunaratne, P.H., Garcia, A.M., Lu, X., Hulyk, S.W., Nanavati,
Kowis, C.R., Sneed, A.J., Martin, R.G., Muzny, D.M., Nanavati,
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (16-NOV-20 Rockville, MD 20850,
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequencing Center
Center code: BCM-HGSC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Baylor College of Medicine Human Genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Pa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NIH-MGC Project URL: http://mgc.nci.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Strausberg, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cDNA Library Preparation: Michael
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BC042531.1 GI:27503509
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                                                                                                                                                                                                                                                                                                                                                       Gibbs,
                                                                 /mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:4837724"
/tissue_type="Testis"
/clone_Tib="NIH_MGC_97"
/lab_host="DH10B"
                                                                                                                                                                           Location/Qualifiers
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                                                                                                                                organism="Homo sapiens"
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mol_type="genomic DNA"
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Pred. No. 1.2;
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RESULT 15
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Matches 24
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Best Local :
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                                                                                                                                         125
AV191114 Yuji Kohara unpublished cDNA:Strain N2 hermaphrodite embryo Caenorhabditis elegans cDNA clone yk590b5 5', mRNA sequence. AV191114
                                                                                                                                                                                                            24;
                                                                                                                                                                          1 CGAACCAGGGCACGTGCAATGGCGA 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted (02-AUG-2001) Asao Fujiyama, The Institute of Physical and Chemical Research (RIKEN), Genomic Sciences Center (GSC); 1-7-22 Suehiro-chou, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan (E-mail:chimpbes@gsc.riken.go.jp/, URL:http://hgp.gsc.riken.go.jp/, Tel:81-45-503-9111, Fax:81-45-503-9170)
Clones are derived from the chimpanzee BAC library RPCI-43 This BAC end was generated during the R&D process and may have higher chance
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Pan troglodytes
Pan troglodytes
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Pan.
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Totoki,Y., Watanabe,H. and Sakaki,Y
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R.Site 2 : EcoRI.
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AUTHORS
Search completed: February 9, 2005, 21:55:45
Job time : 2451.85 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Expressed genes in C.elegans
Unpublished (1999)
Contact: Yuji Kohara
Genome Blology Lab.
National Institute of Genetics
Yata 1111, Mishima, Shizuoka 411, Japan
Tel: 81-559-81-6854
Fax: 81-559-81-6855
Email: ykohara@lab.nig.ac.jp.
Location/Qualifiers
                                                                              303 CGAACCAGGCAGGTGAAATGGAGA 327
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Caenorhabditis elegans
Eukaryota, Metazoa; Nematoda; Chromadorea; Rhabditida;
Ehabditoidea; Rhabditidae; Peloderinae; Caenorhabditis.

1 (bases 1 to 360)
Kohara,Y., Shin-i,T., Thierry-Mieg,J., Thierry-Mieg,D., Mitsuki,H.,
Nishigaki,A., Wotohashi,T., Zeng,Q., Watanabe,H., Sugimoto,A.,
Sano,M., Miyata,A., Mitani,Y., Iida,K., Uesugi,H., Sugiyama,Y. and
Nomoto,H.
                                                                                                          1 CGAACCAGGGCACGTGCAATGGCGA 25
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EST.
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hermaphrodite embryo"
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Maximum Match 100%
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Maximum DB seq length: 2000000000
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  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Gapop 10.0 , Gapext 1.0
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| RESULT 1
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| AX504312 | | • | 100.0 | • | | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | • | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | , |
| | | 147436 | 128993 | 128993 | 98472 | 98472 | 98472 | 9318 | 8630 | 8630 | 8630 | 8630 | 6505 | 4669 | 4669 | 4669 | 4669 | 4669 | 4646 | 4646 | 4646 | 4646 | 4646 | 4646 | 4646 | 4646 | 4646 | |
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| DNA | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
| linear PAT | | AC079303 | AX707913 | AX706983 | AC005068 | AX707905 | AX 706975 | AR028672 | AX012321 | AX012320 | AR306492 | AR306491 | AR028671 | AR405961 | AR363344 | AR203322 | 108557 Sequence 3 | AR091275 | M14758 H | ഥ | | | | AR380622 | 149610 5 | COSTOS | 00004 | 2017195 |
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| RESULT 2 AX504329/c LOCUS DEFINITION ACCESSION VERSION VERYWORDS SOURCE ORGANISM | Qy
da | Query Match
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Matches 2! | AX504312 LOCUS LOCUS DEFINITION ACCESSION VERSION KEYWORDS SOURCE ORGANISM REFERENCE AUTHORS TITLE JOURNAL FEATURES BOUICCE ORIGIN |
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| AX504329 25 bp DNA linear PAT 27-SEP-2002 Sequence 32 from Patent WO0234291. AX504329 AX504329.1 GI:23386141 Homo sapiens (human) | 1 GCTTGTGATCCACGGACACTCCTAC 25 | Query Match 100.0%; Score 25; DB 6; Length 25;
Best Local Similarity 100.0%; Pred. No. 0.04;
Matches 25; Conservative 0; Mismatches 0; Indels 0; Gaps 0; | AX504312 AX504312 AX504312 AX504312 AX504312.1 GI:23386130 Homo sapiens (human) Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 Colgan,S.P. Compositions and methods for treating hematologic malignancies and multiple drug resistance Patent: WO 0234291-A 15 02-MAY-2002; Patent: WO 00234291-A 15 02-MAY-2002; THE BRIGHAM AND WOMEN'S HOSPITAL, INC. (US) Location/Qualifiers 125 //organism="Homo sapiens" //mol_type="unassigned DNA" //db_xref="taxon:9606" |

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J. Biol. Chem. 265 (1), 506-514 (1990)
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Patent: WO 0234291-A 32 02-MAY-2002;
THE BRIGHAM AND WOMEN'S HOSPITAL, INC.
LOCATION/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                    Draft entry and computer-readable by I.B.Roninson, 27-OCT-1989.
                                                                                                                                                                                                                                                                                                                                                                                                                                    Original source text: Human multidrug resistant cell line
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1 (bases 1 to 210)

Chen,C.J., Clark,D., Ueda,K., Pastan,I., Gottesman,M.M. and Roninson,I.B.
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 100.0%; Score 25; DB 9; Length 210; 100.0%; Pred. No. 0.043; tive 0; Mismatches 0; Indels
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RESULT 6
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AX481416/c
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BD171402 3843 bp
Method for predicting side effects
used therefor.
BD171402
BD171402.1 GI:28412692
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I33621
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                                                                                                                                                                                                                                                                                                      Kreutzer,R., Limmer,S., Rost,S. and Hadwiger,P. Method for inhibiting the expression of a target Patent: WO 02055693-A 30 18-JUL-2002; Ribopharma AG (DE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                   AX481416
Sequence 30 from Patent
AX481416
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Bhttnagar, S.K., George, A.L. Jr. and Naz.
Amplification of mucleic acid sequences
Patent: US 5593840-A 1 14-JAN-1997;
                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                     /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
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US 5593840
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WO02055693.
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                                      A linear PAT 18-FEB-2003 immunosuppressant and primer
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Sequence 1 from Patent WO0192877.
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
1 (bases 1 to 3843)
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                                                 25;
                                                                                                                                                                  Sorrentino, B. and Schuetz, J.
Method of identifying and/or isolating stem cells
Patent: WO 0192877-A 1 06-DEC-2001;
ST. JUDE CHILDREN'S RESEARCH HOSPITAL (US)
Location/Qualifiers
                                                                                                                                                                                                                                               Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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GCTTGTGATCCACGGACACTCCTAC 25
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JP 200223769-A/1
13-AUG-2002
31-JAW-2001 JP 2001024723
ICHIRO IEIRI
C12N15/09,C12Q1/68,C12N15/00
Method for predicting side effects of primer used
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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                                                 Score 25; DB 6; Length 3860; Pred. No. 0.047; Mismatches 0; Indels
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RESULT 8 AX322789/c

> RESULT 10 AR452556/c

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AR452556

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PAT 20-FEB-2004

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 2096
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Patent: JP 200252381-A 1 23-JUL-2002;
Wolfgang STREMMEL
OS Homo Saptens
PN JP 2002522381-A/1
PD 23-JUL-2002
PF 06-AUG-1999 JP 2000563262
PR 06-AUG-1999 DE 198 35 526.2,15-DEC-19
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Phosphatidylcholine as a m
BD190394
BD190394.1 GI:33000133
JP 2002522381-A/1.
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Sequence 3 from Patent WO0192877.
XX322789
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Method of identifying and/or isolating stem cells
Patent: WO 0192877-A 3 06-DEC-2001;
ST. JUDE CHILDREN'S RESEARCH HOSPITAL (US)
Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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25; Conserv
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CC
FH Key
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  GCTTGTGATCCACGGACACTCCTAC 2072
              GCTTGTGATCCACGGACACTCCTAC 25
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llarity 100.0%;
Conservative 0
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06-AUG-1998 DE 198 35 526.2,15-DEC-1998
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Pred. No. 0.047;
D; Mismatches
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                                                                                          4192 bp
Homo sapiens P-glycoprotein (mdr1)
AF016535
AF016535.1 GI:2353263
            Chen, C.J.,
                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens (human)
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Sequence 1 from Patent DE19857750.
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Gottesman, M.M. and Roninson, I.B.
                                                            Homo sapiens
                                                                         Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                   Patent: DE 19857750-A 1 24-FEB-2000; STREMMEL WOLFGANG (DE)
                                                                                                                                                                                                                                                                                                                                                                                                           Stremmel, W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Phosphatidylcholine as medication with protective effect large intestinal mucosa Patent: US 667/319-A 1 13-JAN-2004;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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Stremmel, W.
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           (bases 1 to 4192)
len,C.J., Chin,J.E., Ueda,K.,
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           Clark, D.P.,
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/gene="mdr1"
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                                                                      /citation=[1]
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                                                                                               'gene="mdr1"
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2 (bases 1 to 4192)
Chen.G., Duran.G.E., Steger,K.A., Lacayo,N.J., Jaffrezou,J.P.,
Dumontet,C. and Sikic,B.I.
Multidrug-resistant human sarcoma cells with a mutant
P-glycoprotein, altered phenotype, and resistance to cyclosporins
D.B. Biol. Chem. 272 (9), 5974-5982 (1997)
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87028230
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                                SVDGQDĪRTINVRFLREIIGVVSQEPVLFATTIAENIRYGRENVTMDĒIEKĀVKEĀNĀ
YDETIKLEHKEDTLVGERGAQLSGGOKQRIA TARALVENBEI LILLDEATSALDTESSA
VUQVALDKARKGRTI VIAHKLSTVENADVJAGEDDGVIVEKGHNDELMKEKGI YEKL
VUMQTAGNEVELENAADESKS EIDALEMSSNDSRSSLIRKRSTRRSVRGSQAQDRKLS
TKEALDESI EPVSFWRIMKLMITEMPY FVVQVPCAI INGGLQPAFA I IFSKI IGVETR
IDDPETKRQNSNLFSLLFLALGI I SEI TFFLQGFTFGKAGEI LTKRLHXWVFERSMLRQ
DVSWFDDPKNTTGALTTRLANDAQVKGAI GSSLAVITQNI ANLGTGI I IS FI YGWQL
TLLLLAI VPI IA IAGVVEMKNLSGQALKDKKELEGAGKIA TEAA LENFRTVSLTQEQK
FEHMYAQSLQVP YRNSLRKAHI FGITFS FTQAMMYFS YAGCFRFGAYLVAHKLMSFED
VLLVESAVVEGAMAVGQVSSFAPDYAKAKI SAAHI IMI IEKTPL LDYSTEGLMENTL
EGNVTFGEVVFNYPTREDI PVLQGISLEVKKQQTLALVGSGCGKSTVQOLLERFYDP
LAGKVLLDGKEI KRLNVQMLRAHLGI VSQEP IL FDCS I AENI AYGDNSRVVSQEEI VR
AAKEANIH PI ESLENKYSTRYGDKGTOLSGGOKQR IA IARALVRQPHILLLDEATSA
LDTESEKVVQEALDKAREGRTCI VI AHRLSTI QNADLI VVFQNGRVKEHGTHQQLLAQ
LDTESEKVQEALDKAREGRTCI VI AHRLSTI QNADLI VVFQNGRVKEHGTHQQLLAQ
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FFMNLEEDMTRYAYYSGIGAGVLVAAYIQVSFWLAAAGRQIHKIRKQFFHAIMRQEI
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FRNVHFSYPSRKEVKILKGLNLKVQSGQTVALVGNSGCGKSTTVQLMQRLYDFTEGMV
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Sequence 5 from patent US 5830697.
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Sikic, B.I. and Chen, G.
P-glycoprotein mutant resistant to cyclosporin modulation
Patent: US 5830697-A 1 03-NOV-1998;
Location/Qualifiers
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Sikic, B.I. and Chen,G.
P-glycoprotein mutant resistant to cyclosporin modulation
Patent: US 5830697-A 5 03-NOV-1998;
Location/Qualifiers
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completed: February
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                                                                                                                                                                                                                                                                                         Patent: JP 1990100680-A 1
SUNTORY LTD
OS Homo sapiens
PN JP 1990100680-A/1
PD 12-APR-1990
PF 05-OCT-1988 JP 19882:
PI UEDA KAZUMITSU, KOMAI
PC C12N15/12,C12N1/21,CC
CC strandedness: Single
CC topology: Linear;
CC *source: tissue_type:
CC *source: clone=SAM11.
FH Key
FT CDS 138.
FT CDS 138.
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E02326
E02326.1 GI:2170561
JP 1990100680-A/1.
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                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 4378)

Ueda, K. and Komano, T.

HUMAN NORMAL CELL-DERIVED MDR RELATED GENE

Patent: JP 1990100680-A 1 12-APR-1990;
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JP 1990100680-A/1
12-APR-1990
05-OCT-1988 JP 1988251475
UEDA KAZUMITSU, KOMANO TORU
C12N15/12,C12N1/21,C12Q1/68;
strandedness: Single;
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                                                                                                         Conservative
                                                                                                                                                                         /organism="Homo sapiens"
/mol_type="genomic RNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                               Location/Qualifiers
                                                                                                        100.0%; Score 25; DB 6;
100.0%; Pred. No. 0.047;
vative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                              tissue_type=Adrenal gland; clone=SAM1132;
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/gene='MDRA1'
3981. .>4378.
               2005,
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relating
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Maximum Match 100%
Listing first 45 summaries
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  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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                    US-08-461-823-1
US-08-784-649A-5
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US-09-949-016-14353
US-09-252-991A-3469
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| 64.0 | 64.8 | | 64.8 | 64.8 | 64.8 | 64.8 | 64.8 | 64.8 | 64.8 | 64.8 | 64.8 | 64.8 | 64.8 | 66.4 | 66.4 | 66.4 |
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| US-09-949-016-26189 | US-09-949-016-16688 | US-09-949-016-15043 | US-09-949-016-15042 | US-09-949-016-13903 | US-09-949-016-16897 | US-09-949-016-12820 | US-09-252-991A-11533 | US-09-252-991A-11651 | US-09-252-991A-11783 | US-09-252-991A-11705 | US-09-949-016-174876 | US-09-949-016-174875 | US-09-252-991A-11850 | US-09-949-016-17556 | US-09-949-016-13429 | US-09-949-016-205912 |
| Sequence 26189, i | 16 | Sequence 15043, | Sequence 15042, | Sequence 13903, | Sequence 16897, | Sequence 12820, | Sequence 11533, | Sequence 11651, | Sequence 11783, | Sequence 11705, | Sequence 174876, | Sequence 174875, | Sequence 11850, | Sequence 1/556, | Sequence 13429, | Sequence 205912, |

ALIGNMENTS

| ZEELEN TOLFFRARS BUNCHARGO | RESULT 1 US-08-461-823-1/c US-08-461-823-1/c (Sequence 1, Application US/08461823) Patent No. 5593840 ; GENERAL INFORMATION: APPLICANT: Bhatnagar, Satish K. APPLICANT: George Jr., Albert L. APPLICANT: Nazarenko, Irina TITLE OF INVENTION: AMPLIFICATION OF NUCLEIC ACID SEQUENCES: NUMBER OF SEQUENCES: 27 CORRESPONDENCE ADDRESS: ADDRESSEE: OncorPharm, Inc. STREET: 200 Perry Parkway CITY: Gatthersburg STATE: Maryland COUNTRY: USA ZID. 20877 |
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Query Match Best Local

Similarity

100.0%; Score 25; DB 1; Length 2726; 100.0%; Pred. No. 0.019;

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; LENGTH: 3988
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-762-195-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: 34691/208520
CURRENT APPLICATION NUMBER: US/09/762,195
CURRENT FILING DATE: 2001-02-05
PRIOR APPLICATION NUMBER: PCT/EP99702426
PRIOR FILING DATE: 1999-08-06
PRIOR APPLICATION NUMBER: 198 35 526 2 DE
PRIOR FILING DATE: 1998-08-06
PRIOR APPLICATION NUMBER: 198 57 570.8 DE
PRIOR APPLICATION NUMBER: 198 57 570.8 DE
PRIOR APPLICATION NUMBER: 198 57 570.8 DE
PRIOR FILING DATE: 1998-12-15
NUMBER OF SEQ ID NOS: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 1, Application US/08784649A Patent No. 5830697
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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Patent No. 6677319
TENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Stremmel, Wolfgang
IITLE OF INVENTION: Phosphatidylcholine as Medication with
IITLE OF INVENTION: Protective Effect on Large Intestinal Mucosa
ATTORNEY/AGENT INFORMATION:

NAME: Sherwood, Pamela J

REGISTRATION NUMBER: REG.NO. 5830697 36,677

REFERENCE/DOCKET NUMBER: 06037/007001
                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
STREET: 2200 Sand Hill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Sikic, Branimir I
APPLICANT: Chen, Gang
TITLE OF INVENTION: P-GLYCOPROTEIN MUTANT RESISTANT TO
TITLE OF INVENTION: CYCLOSPORIN MODULATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                         APPLICATION NUMBER: US/08/784,649A FILING DATE: CLASSIFICATION: 435
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STATE: CA
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1 Similarity 100.0%; Pred. No. 0.02;
25; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent No. 5830697
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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Best Local &
                                                               Matches
                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 415-854-0875
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                          TELEFAX: 415-854-0875
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
ODERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/784,649A
FILING DATE:
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
STREET: 2200 Sand Hill Road
                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Sherwood, Pamela J
REGISTRATION NUMBER: REG.NO.
REFERENCE/DOCKET NUMBER: 060
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Sikic, Bran
APPLICANT: Chen, Gang
TITLE OF INVENTION: P-
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                                                                                                                                            TOPOLOGY: 1
MOLECULE TYPE:
                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: CDNA
                                                                                                                                                         STRANDEDNESS: single TOPOLOGY: linear
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                       TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Menlo Park
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2168
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                               1 GCTTGTGATCCACGGACACTCCTAC 25
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GCTTGTGATCCACGGACACTCCTAC 2144
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415-854-0875
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                                                               Conservative
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100.0%; Pred. No. 0.
ative 0; Mismatches
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                                                                            100.0%;
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                                                             Score 25; DE
Pred. No. 0.0
0; Mismatches
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US-08-181-471-2/c
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; Sequence 1167, Application US/09023655,
; Patent No. 6607879
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US-08-181-471-2
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Best Local Similarity
Matches 25; Conserv
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APPLICANT: Lishkc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Lishko, Valeryi K.
TITLE OF INVENTION: COMPOSITIONS TO HAIR FOLLICLES
                                                         APPLICANT: Susan G. Stuart
APPLICANT: Susan G. Stuart
APPLICANT: Jeffrey J. Seilhamer
APPLICANT: OMPOSITION
COMPOSITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: CI
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                           TITLE OF INVENTION: COTITLE OF INVENTION: EX NUMBER OF SEQUENCES: 1
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ZIP: 921
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STREET: 12526 High Bluff Drive, Suite 300
                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                  NAME/KEY:
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               ADDRESSEE:
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                                                                                                                     INFORMATION:
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3174 PORTER DRIVE
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                                                                     THE DETECTION OF BLOOD CELL GENE
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US-08-752-447-1/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (650) 845-4166
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 4646 base pairs
                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Mechetner
APPLICANT: Roninson,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Zeller, Karen J.
REGISTRATION NUMBER: 37,071
REFERENCE/DOCKET NUMBER: PA
TELECOMMUNICATION INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS 6.2
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/023,655
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IMMEDIATE SOURCE:
LIBRARY: GENER
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SOPTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/752,447
FILLING DATE: 15-NOV-1996
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                        ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
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Local Similarity 100.0%; Pred. No. 0.
hes 25; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
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CLASSIFICATION:
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                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                              CITY: Chicago
STATE: Illinois
                                                                                                                                                                                                                                   STREET:
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300 South Wacker Drive, Seventh Floor
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Methods and Reagents for Preparing
Using Immunoligcal Agents Specific
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                                                                            Version
                                                                            #1.30
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for P-glycoprotein
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ATTORNEY/AGENT INFORMATION:

NAME: NO. 5994088nan, Kev REGISTRATION NUMBER: 35,3 REFERENCE/DOCKET NUMBER:

95,1121

TELEPHONE: 312-913-0001

312-913-9808

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                                                                                                                     CLASSIFICATION:
PRIOR APPLICATION NUMBER: 08/752,447
APPLICATION NUMBER: 08/752,447
EILING DATE: 15-NOV-1996
ATTORNEY/AGENT INFORMATION:
NAME: NO. 6365357nan, Kevin E
REGISTRATION NUMBER: 35,303
REFERENCE/DOCKET NUMBER: 95,1121
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-316-167-1/c
                                                                            TELEFAX: 312-913-9808 INFORMATION FOR SEQ ID NO:
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INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                              SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RITLE OF INVENTION: Methods and Reagents for Preparing and Presents of Invention: Using Immunoligical Agents Specific for P-glycoprotein
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LOCATION:
TYPE: nucleic ac
STRANDEDNESS: BI
TOPOLOGY: linear
                                                                                                                                                                                                                                                                  FILING DATE:
                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
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LOCATION:
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                                             LENGTH:
                                                                                                          TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 4669 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPOLOGY:
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25; Conservative
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                             4669 base pairs ucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                              USA
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Roninson, Igor B
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1..424
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4265..4669
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Pred. No. 0.02;
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Query Match
Best Local Similarity
Watches 25; Conserva
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US-09-397-233-1/c
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Patent No. 6630327
GENERAL INFORMATION:
 Query Match
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INFORMATION FOR SEQ ID NO: 1:
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LOCATION:
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                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/09/397,233
FILING DATE: 16-Sep-1999
CLASSIFICATION: «Unknown»
ATTORNEY/AGENT INFORMATION:
NAME: NO. 6630327nan, Kevin E
REGISTRATION NUMBER: 35,303
REFERENCE/DOCKET NUMBER: 95,1121-0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                LOCATION: 4265..4669
SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                   FEATURE:
                                                                                                                                                  FEATURE:
                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Roninson, Igor B
TITLE OF INVENTION: Methods and Reagents for Preparing and
Using Immunological Agents Specific for P-glycoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2454 GCTTGTGATCCACGGACACTCCTAC 2430
                                                                                                                                                                                                  FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
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Roninson, Igor B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
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                                                                                  NAME/KEY:
                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
ZIP: 60606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Chicago
STATE: Illinois
                                                                                                                                    NAME/KEY:
                                                                                                                                                                 LOCATION:
                                                                                                                                                                                  NAME/KEY:
                                                                                                                                                                                                                                   TOPOLOGY:
                                                                                                                                                                                                                                                                                 LENGTH: 4669 base pairs
                                                                                                                                                                                                                                                                                                                                               TELEPHONE: 312-913-0001
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STREET: 300 South Wacker Drive
                                                                                                               OCATION:
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4265..4669
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1..424
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425..4264
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1..424
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100.0%; Pred. No. 0.0
ative 0; Mismatches
100.0%; Score 25;
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 DB 4;
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Length 4669;
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RESULT 10
$206352-3/c
;Patent No. 5206352

;Patent No. 5206352

;Michael M.

TITLE OF INVENTION: COMPOSITIONS FOR CLONES CONTAINING DNA
;SEQUENCES ASSOCIATED WITH MULTIDRUG RESISTANCE IN HUMAN CELLS
;NUMBER OF SEQUENCES: 4
;CURRENT APPLICATION DATA;
;CURRENT APPLICATION DATA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 5206352-3/c
; Patent No. 5206352
; Patent No. 5206352
; APPLICANT: Roninson, Igor B.; Pastan Ira H.; Gottesman,
; Michael M.
; MICHAEL M.
; TITLE OF INVENTION: COMPOSITIONS FOR CLONES CONTAINING DNA
; SEQUENCES ASSOCIATED WITH MULTIDRUG RESISTANCE IN HUMAN CELLS
; NUMBER OF SEQUENCES: 4
; CURRENT APPLICATION DATA;
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US-08-793-610-5/c
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Best Local Similarity
Matches 25; Conserv
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Matches
                                           Sequence 5, Application US/08793610
Patent No. 5858744
                                                                                                                                                                                                                                             Query Match
                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/
FILING DATE: 24-SEP-1990
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/
FILING DATE: 24-SEP-1990
PRIOR APPLICATION DATA:
APPLICANT:
                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 892,575
FILING DATE: 01-AUG-1986
APPLICATION NUMBER: 845,610
FILING DATE: 28-MAR-1986
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FILING DATE: 01-AUG-1986
APPLICATION NUMBER: 845,610
FILING DATE: 28-MAR-1986
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 BAUM, Christopher
STOCKING-HARBERS, Carol
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tive 0; Mismatches
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100.0%; Pred. No. 0.02;
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RESULT 13
US-09-306-417-1/c
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PATENTIAN PC-DOS/MS-DOS

OPERATING SYSTEM: PC-DOS/MS-DOS

OPERATING SYSTEM: PC-DOS/MS-DOS

OURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/793,610

FILING DATE: 07-MAR-1997

PRIOR APPLICATION DATA:

APPLICATION NUMBER: DE P 44 31 973.8

FILING DATE: 08-SEP-1994

PRIOR APPLICATION DATA:

APPLICATION NUMBER: DE 195 03 952.1

FILING DATE: 07-FEB-1995

PRIOR APPLICATION DATA:

APPLICATION NUMBER: PCT/EP95/03175

PRIOR APPLICATION DATA:

APPLICATION NUMBER: PCT/EP95/03175

PRIOR APPLICATION INTOMATA:

APPLICATION DATA:

APPLICATION STATE: 10-AUG-1995

ATTONNEY/AGENT INFORMATION:

NAME: Berman, Richard J.
                                                                                                                 EARLIER FILING DATE: 1998-05-08
NUMBER OF SEQ ID NOS: 17
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 1
                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Heinrich-Pette-Institut
APPLICANT: INVENTION: Retroviral Gene Transfer Vectors
TITLE OF INVENTION: Retroviral Gene Transfer Vectors
FILE REFERENCE: P50491
                                                                                                                                                                                                                                                                                                                                                                            Sequence 1, Application US/09306417 Patent No. 6548301
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Best Local Similarity
                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/306,417
CURRENT FILING DATE: 1999-05-06
EARLIER APPLICATION NUMBER: DE 198 22 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (202) 638-481
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nikaido, Marmelstein, Murray & Oram LLP
STREET: 655 Fifteenth Street N.W. Suite 330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
OTHER INFORMATION: Description of Artificial Sequence: proviral
                        ORGANISM: Artificial Sequence FEATURE:
                                                                       TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: (202)638-5000
                                                                                             ENGTH: 8630
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Berman, Richard J.
REGISTRATION NUMBER: 39,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (202) 638-4810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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OTHER INFORMATION: plasmid

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                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 2
LENGTH: 8630
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/306,417
CURRENT FILING DATE: 1999-05-06
EARLIER APPLICATION NUMBER: DE 198 22 115
EARLIER FILING DATE: 1998-05-08
NUMBER OF SEQ ID NOS: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Heinrich-Pette-Institut
TITLE OF INVENTION: Retroviral Gene Transfer Vectors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            equence 2, Application US/09306417 atent No. 6548301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: P50491
NAME/KEY: misc feature LOCATION: (161)..(677)
OTHER INFORMATION: 5'-LTR
                                                                                FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)..(160)
OTHER INFORMATION: plasmid
                                                                                                                                                         NAME/KEY: misc_feature
LOCATION: (1)..(8630)
OTHER INFORMATION: retroviral expression vector SFbeta91mSA1
                                                                                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc_feature
LOCATION: (1).. (8630)
OTHER INFORMATION: retroviral expression vector SFbeta71m4
                                                                                                                                                                                                                                                      OTHER INFORMATION: Description of Artificial Sequence: proviral OTHER INFORMATION: plasmid DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: mat_peptide
LOCATION: (1220)..(5062)
OTHER_INFORMATION: m4 mdr-1 cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: misc feature
LOCATION: (161)...(677)
OTHER INFORMATION: 5'-LTR
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LOCATION: (1)...(160)
OTHER _INFORMATION: plasmid backbone (pUC)
                                                                                                                                                                                                                                                                                                    FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: (5775)..(8630)
OTHER INFORMATION: plasmid backbone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LOCATION: (5215)..(5774)
OTHER INFORMATION: 3'-LTR
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LOCATION: (5775)..(863
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JOCATION: (532)..(1219)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
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                                                                                    backbone (pUC)
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TELEPAX: (202)000 6:
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 9318 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
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Best Local S
Matches 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 6,
                                                                                                                                                                                                                                                                                                  FILING DATE: 07-MAR-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE P
FILING DATE: 08-SEP-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE 195
FILING DATE: 07-FEB-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
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LOCATION: (5775)..(8630)
OTHER INFORMATION: plasmid backbone (pUC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: misc_feature
LOCATION: (5215)..(5774)
OTHER INFORMATION: 3'-LTR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
APPLICANT:
APPLICANT:
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OTHER INFORMATION: mSA1 mdr1 cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: mat_peptide
LOCATION: (1220)..(50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: 5'UTR
LOCATION: (532)..(1219)
                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95/03175
FILING DATE: 10-AUG-1995
ATTORNEY/AGENT INFORMATION:
NAME: 10-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/793,610
                                                                                                                                         TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: D
                                                                                                                                                                 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                              NAME: Berman, Richard J.
REGISTRATION NUMBER: 39,105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET:
                                                                                                                        TELEPHONE:
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5858744
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3: Nikaido, Marmelstein, Murray & Oram LLP
655 Fifteenth Street N.W. Suite 330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BAUM, Christopher
STOCKING-HARBERS, Carol
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 25; DB 4; Length 8630; 100.0%; Pred. No. 0.022; ative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Post-processing: Minimum Match 0%
Maximum Match 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Minimum DB seq length: 0
Maximum DB seq length: 200000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Perfect score:
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                                                                                                                                                                                                                                                                                                        Published_Applications_NA:*

1: /cgn2_6/ptcdata/1/pubpna/US07_PUBCOMB.seq:*

2: /cgn2_6/ptcdata/1/pubpna/US06_NEW_PUB.seq:*

3: /cgn2_6/ptcdata/1/pubpna/US06_NEW_PUB.seq:*

4: /cgn2_6/ptcdata/1/pubpna/US06_PUBCOMB.seq:*

5: /cgn2_6/ptcdata/1/pubpna/US07_NEW_PUB.seq:*

6: /cgn2_6/ptcdata/1/pubpna/US08_NEW_PUB.seq:*

7: /cgn2_6/ptcdata/1/pubpna/US08_NEW_PUB.seq:*

8: /cgn2_6/ptcdata/1/pubpna/US08_PUBCOMB.seq:*

9: /cgn2_6/ptcdata/1/pubpna/US08_PUBCOMB.seq:*

9: /cgn2_6/ptcdata/1/pubpna/US08_PUBCOMB.seq:*
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25
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Copyright (c) 1993 - 2005 Compugen Ltd.
/ cgn2_6/ptodata/1/pubpna/US09_BPUBCOMB.seq:*
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/cgn2_6/ptodata/1/pubpna/US09_EW_PUBCOMB.seq:*
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Pred. No. re greater t is derived is the number of results predicted by chance to have a ster than or equal to the score of the result being printed. à analysis of the total score distribution

SUMMARIES

| 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 | Result |
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| ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ | Score |
| 100.0
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Match |
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| 18
99
113
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17 | B |
| US-10-384-339C-30 US-09-866-866A-1 US-09-866-866A-3 US-09-856-020-30 US-10-072-621-2 US-10-097-340-1 US-10-097-926A-258 US-09-968-007A-747 US-10-641-643-1167 US-10-343-657-1 | ID |
| Sequence 30, Appli
Sequence 1, Appli
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| 45 | 44 | 43 | 42 | 41 | 40 | 39 | 8 | . U. | 9 6 | ω
G | 34 | 33 | 32 | 31 | 30 | 29 | 28 | 27 | 26 | 25 | 24 | 23 | 22 | 21 | 20 | 19 | 18 | 17 | 16 | 15 | 14 | 13 | 12 |
| 17 | 17 | 17 | 17 | 17 | 17 | 17 | 1.7 | ; <u>-</u> | | 17 | 17 | 17 | 17 | 17 | 17.2 | 17.2 | 7. | 17.2 | | 17.2 | | -1 | • | 17.2 | 17.6 | 19.6 | 24 | 24 | 24 | 25 | 25 | 25 | 25 |
| 68.0 | 68.0 | 68.0 | 68.0 | 68.0 | α. | | α | œ. | | . « | . 8 | 8 | 68.0 | 68.0 | | ٠ | ~ | 68.8 | m | 68.8 | m | 68.8 | ന | 68.8 | õ | 78.4 | 96.0 | 96.0 | 96.0 | 100.0 | 100.0 | 100.0 | 100.0 |
| 1929 | 1781 | 1776 | 1535 | 1535 | 1464 | 1198 | . 942 | 200 | 100 | 550 | 550 | 463 | 463 | 388 | 136328 | 26555 | 26555 | 2390 | 2390 | 1005 | 665 | 665 | 625 | 625 | 334 | | 4. | | 3852 | 8630 | 8630 | 4669 | 4646 |
| <u>, , , , , , , , , , , , , , , , , , , </u> | Η. | ۳. | μ | <u>_</u> | | 17 | | : : | | ٠. | . 1 | 17 | <u></u> | 18 | 15 | 17 | 9 | 17 | 13 | 17 | 17 | 13 | 17 | 13 | 18 | 9 | 17 | 17 | 15 | φ | 9 | 18 | 18 |
| US-10-282-122A-29808 | US-10-425- | US-10-425-114- | US-10-027-632- | US-10-02/-632-265/0 | - 665- 525 OT - 511 | US-10-369- | 007-4771-707-01-50 | 00-03-03E0-03E-00-00 | _ | | 8 | US-10-027-632-70301 | US-10-027-632-70301 | -10-425-115 | -10-101-510-127 | US-10-227-646-161 | US-09-860-670-161 | -632 | -10-027-632-10177 | -10-369-493-26051 | 0-027-632- | -10-027-632-12698 | US-10-027-632-290365 | 632-29036 | US-10-425-115-130740 | 74-269 | -359A- | US-10-619-359A-1 | US-10-101-433A-1 | US-09-306-417-2 | - 1 | US-10-680-516-1 | US-10-775-169-198 |
| Sequence 29808, A | | 13600, | e 200700, | 1000 | 365708 | 30343 | 20224 | 28884 | Semience 1088. An | 1088 21 | | 70301, | 70301, | 38049, | 12/, AP | 161, | 161, / | e 101 | Sequence 101776, | | 126989, | 12698 | | | | 2 | , e | | ۲ | | , - | . ₽ | Sequence 198, App |
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ALIGNMENTS

US-10-384-339C-30/c

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US-10-384-339C-30
                                                                                                                                                                                                                                             ; SOFTWARE: PatentIn Ver. 2; SEQ ID NO 30; LENGTH: 3840; TYPE: DNA; ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/384,339C
CURRENT FILING DATE: 2003-03-07
PRIOR APPLICATION NUMBER: PCT/EP02/00152
PRIOR PILING DATE: 2002-01-09
PRIOR APPLICATION NUMBER: DE 1010586.5
PRIOR PILING DATE: 2001-01-09
PRIOR PILING DATE: 2001-01-09
PRIOR APPLICATION NUMBER: DE 10155280.7
PRIOR FILING DATE: 2001-10-26
PRIOR FILING DATE: 2001-11-29
PRIOR APPLICATION NUMBER: DE 10158411.3
PRIOR PILING DATE: 2001-11-29
PRIOR APPLICATION NUMBER: DE 10160151.4
PRIOR PILING DATE: 2001-12-07
NUMBER OF SEQ ID NOS: 173
SOFTWARE: PALENTIN Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 30, Application US/10384339C Publication No. US20040175703A1 GENERAL INFORWATION:
Query Match 100.0%; Score 25; DB 18; Best Local Similarity 100.0%; Pred. No. 0.027; Matches 25; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Kreutzer, Roland
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR INHIBITING EXPRESSION OF A TARGET GE
FILE REFERENCE: 20200/2002
                                                                                                                                                         PUBLICATION INFORMATION:
TITLE: mdr-1
PATENT DOCUMENT NUMBER: AF016535
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                                                                 Length 3840;
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US-09-866-866A-3/c
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-866A-1
                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/866,866A
CURRENT FILING DATE: 2001-08-30
PRIOR APPLICATION NUMBER: 09/584,586
PRIOR FILING DATE: 2000-05-31
PRIOR APPLICATION NUMBER: PCT/US99/11825
PRIOR APPLICATION NUMBER: CT/US99/11825
PRIOR FILING DATE: 1999-05-27
PRIOR PPLICATION NUMBER: 60/086,988
PRIOR FILING DATE: 1998-05-28
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US-09-866-866A-1/c
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PRIOR FILING DATE: 2000-05-31
PRIOR PELICATION NUMBER: PCT/US99/11825
PRIOR FILING DATE: 1999-05-27
PRIOR PAPPLICATION NUMBER: 60/086,988
PRIOR PILING DATE: 1998-05-28
NUMBER OF SEQ ID NOS: 27
SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.0 SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Sorrentino, Brian
APPLICANT: Schuetz, John
TITLE OF INVENTION: A Method of Identifying and/or Isolating Stem Cells
FILE REFERENCE: 1340-1-021CIP2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 1, Application US/09866866A Patent No. US20020102244A1
GENERAL INFORMATION:
                                                                                        Matches
                                                                                                           Query Match
Best Local Similarity
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Best Local
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APPLICANT: Schuetz, John
TITLE OF INVENTION: A Method of Identifying and/or Isolating Stem Cells
FILE REFERENCE: 1340-1-021CIP2
                                                                                                                                                                                             LENGTH: 3860
TYPE: DNA
ORGANISM: homo sapiens
  2030
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Similarity 100.0%; Pred. No. 0.027;
25; Conservative 0; Mismatches 0
                                   GCTTGTGATCCACGGACACTCCTAC 25
  GCTTGTGATCCACGGACACTCCTAC 2006
                                                                                        Conservative
                                                                                 100.0%; Score 25; DB 9;
100.0%; Pred. No. 0.027;
tive 0; Mismatches 0
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                                                                                    Gaps
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; ORGANISM: Homo sapiens
US-10-072-621-2
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                                                                                                                                                                              US-10-097-340-1/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Reiner, Peter B.
APPLICANT: Connop, Bruce P.
APPLICANT: Connop, Michelle
TITLE OF INVENTION: REGULATION OF AMYLOID PRE
TITLE OF INVENTION: BY MODIFICATION OF ABC T
FILE REFERENCE: 100103.402
CURRENT APPLICATION UNMBER: US/10/072,621
CURRENT FILING DATE: 2002-02-08
NUMBER OF SEQ ID NOS: 10
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US-09-805-020-30/c
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                                                                            Sequence 1, Application US/10097340
Publication No. US20030087250A1
GENERAL INFORMATION:
APPLICANT: John MONAHAN
APPLICANT: Manjula GANNAVARAPU
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Publication No. US20020086384A1
GENERAL INFORMATION:
APPLICANT: LEVINE, Zurit
TITLE OF INVENTION: SPLICE VARIANTS OF ONCOGENES
FILE REFERENCE: 2786-0168P
CURRENT APPLICATION NUMBER: US/09/805,020
CURRENT FILING DATE: 2001-03-13
NUMBER OF SEQ ID NOS: 72
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SEQ ID NO 2
LENGTH: 4643
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SEQ ID NO 30
LENGTH: 4533
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Best Local (
APPLICANT:
                                         APPLICANT:
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NAME/KEY: misc_feature
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                    Sebastian HOERSCH
Shubhangi KAMATKAR
Steve G. KOVATS
                                                                                                                                                                                                                                                                                                                                           Conservative
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Michael MORRISEY

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US-10-007-926A-258/c
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; ORGANISM: Homo Bapiens
US-10-097-340-1
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Best Local 9
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                                                                                                                       PRIOR APPLICATION NUMBER: 60/254,090
PRIOR FILING DATE: 2000-12-08
NUMBER OF SEQ ID NOS: 468
SOPTWARE: PATENTIN Ver. 2.1
SEQ ID NO 258
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APPLICANT:
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                                                                                                                                                                                                                                 TITLE OF INVENTION: GENE EXPRESSION PROFILING OF PRIMARY BREAST CARCINOMAS TITLE OF INVENTION: USING ARRAYS OF CANDIDATE GENES FILE REFERENCE: 1546-R-00 CURRENT APPLICATION NUMBER: US/10/007,926A CURRENT FILING DATE: 2001-12-07 CURRENT FILING DATE: 2001-12-07
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PRIOR FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 60/276,026
PRIOR APPLICATION NUMBER: 60/276,026
PRIOR APPLICATION NUMBER: 60/276,026
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CURRENT FILING DATE: 2002-03-14
PRIOR APPLICATION NUMBER: 60/276,025
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-14
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                                        LENGTH: 4643
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
THER INFORMATION: atp-binding cassette, sub-family bother Information: (mdr/tap), member 1 (ABCB1) gene.
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FILING DATE: 2001/09/26
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BIRNBAUM, DANIEL
NGUYEN, CATHERINE
VIENS, PATRICE
FERT, VINCENT
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Robert C. BAST, Jr.
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o. US20030143539A1
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Pred. No. 0.027;
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Best Local
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SEQ ID NO 459
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity Matches 25; Conserv
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TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Usining Signa TITLE OF INVENTION: Gene Sets

FILE REFERENCE: 689290-71

CURRENT FILING DATE: 2001-10-02

PRIOR APPLICATION NUMBER: US/60/237,172

PRIOR FILING DATE: 2000-10-02

PRIOR APPLICATION NUMBER: US/60/237,173

PRIOR APPLICATION NUMBER: US/60/237,278

PRIOR APPLICATION NUMBER: US/60/237,278

PRIOR APPLICATION NUMBER: US/60/237,278

PRIOR APPLICATION NUMBER: US/60/237,278

PRIOR PILING DATE: 2000-10-02

PRIOR APPLICATION NUMBER: US/60/237,294

PRIOR APPLICATION NUMBER: US/60/237,294

PRIOR APPLICATION NUMBER: US/60/237,295
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PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 1001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US/60/237,278
PRIOR FILING DATE: 2000-10-02
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TYPE: DNA
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5. US20040115625A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
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100.0%; Pred. No. 0.027;
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Pred. No. 0.027;
Mismatches
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PRIOR FILING DATE: 2000-10-02 NUMBER OF SEQ ID NOS: 1001 SOFTWARE: PatentIn version 3.0

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; SEQ ID NO 747
LENGTH: 4646
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-968-007A-747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-10-641-643-1167/c
Sequence 1167, Application US/10641643
Publication No. US20040077003A1
GENERAL INFORMATION:
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                                                                                                                                    US-10-641-643-1167
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Matches
                                                           Matches
                                                                           Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 1167:
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                                                                                                                                                                                                                                                                                                                                                                NAME: Zeller, Karen J.
REGISTRATION NUMBER: 37,071
REFERENCE/DOCKET NUMBER: PA-0001 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: <Unknown>
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS 6.2
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/641,643
FILING DATE: 14-Aug-2003
                                                                                                                                                      CLONE: g187468
SEQUENCE DESCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Susan G. Stuart
Jeffrey J. Seilhamer
TITLE OF INVENTION: COMPOSITION FOR THE DETECTION OF BLOOD CELL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2454 GCTTGTGATCCACGGACACTCCTAC 2430
                                                                                                                                                                                                                IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 1508 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Cocks, Benjamin G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 GCTTGTGATCCACGGACACTCCTAC 25
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GCTTGTGATCCACGGACACTCCTAC 25
                                                                                                                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: INCYTE PHARMACEUTICALS, INC. STREET: 3174 PORTER DRIVE
                                                                                                                                                                                              LIBRARY: GENBANK
                                                                                                                                                                                                                                                                                                                                                   TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: PALO ALTO
                                                                                                                                                                                                                                                                                           LENGTH: 4646 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 25; DB 11 llarity 100.0%; Pred. No. 0.027; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CALIFORNIA
                                                                                                                                                                                                                                                                                                                                               E: (650) 855-05
(650) 845-4166
                                                                           100.0%; Score 25; DB 17; Length 4646; 100.0%; Pred. No. 0.027;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENE EXPRESSION
                                                                                                                                                        SEQ ID NO: 1167 :
                                                           0
                                                           Mismatches
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RESULT 13
US-10-680-516-1/c
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US-10-775-169-198/c
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                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.2
SEQ ID NO 198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 198, Application No. US200 GENERAL INFORMATION:
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                                                                                                                                                     Matches
                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                    APPLICANT: Twine, Natalie
APPLICANT: Dorner, Andrew
APPLICANT: Trepicchio, William
APPLICANT: Trepicchio, William
TITLE OF INVENTION: Method for Monitoring Drug Activities In Vivo
FILE REFERENCE: AMIO1080 (031896-013000)
CURRENT APPLICATION NUMBER: US/10/775,169
CURRENT FILING DATE: 2004-02-11
NUMBER OF SEC ID NOS: 5278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/222,313
PRIOR FILING DATE: 2000-08-01
NUMBER OF SEQ ID NOS: 18
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Mutations of the MDR1 P-glycoprotein that Improve Its TITLE OF INVENTION: Ability to Confer Resistance to Chemotherapeutic Drugs FILE REFERENCE: 00,616-A CURRENT APPLICATION NUMBER: US/10/343,657 CURRENT FILING DATE: 2003-10-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Wyeth
APPLICANT: Burczynski, Michael
APPLICANT: Twine, Natalie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Roninson, Igor B. APPLICANT: Ruth, Adam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
NAME/KEY: CDS
LOCATION: (425)..(4264)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 4646
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                     LENGTH: 4646
                                                                             2454 GCTTGTGATCCACGGACACTCCTAC 2430
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                                                                                                                GCTTGTGATCCACGGACACTCCTAC 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Application US/10775169 o. US20040175743A1
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                                                                                                                                                     Conservative
                                                                                                                                                                    100.0%; Score 25;
100.0%; Pred. No.
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                                                                                                                                                     0; Mismatches
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Pred. No. 0.027;
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                                                                                                                                                                                            DB 18;
                                                                                                                                                                        .027;
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                                                                                                                                                                                          Length 4646;
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                                                                                                                                                     Gaps
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; Sequence 1, Application US/10680516; Publication No. US20040166110A1; GENERAL INFORMATION:

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RESULT 14
US-09-306-417-1/c
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GENERAL INFORMATION:
APPLICANT: Heinrich-Pette-Institut
TITLE OF INVENTION: Retroviral Gene Transfer Vectors
FILE REFERENCE: P50491
CURRENT APPLICATION NUMBER: US/09/306,417
CURRENT FILING DATE: 1999-05-06
EARLIER APPLICATION NUMBER: DE 198 22 115
EARLIER FILING DATE: 1998-05-08
NUMBER OF SEQ ID NOS: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                  Sequence 1, Application US/09306417 Patent No. US20020103144A1
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APPLICATION NUMBER: US/10/680,516

PILING DATE: 07-Oct-2003

CLASSIFICATION: 435

PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Noonan, Kevin E
REGISTRATION NUMBER: 35,303
REFERENCE/DOCKET NUMBER: 95
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-913-0001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Roninson, Igor B
TITLE OF INVENTION: Methods and Reagents for Preparing and
Using Immunoligcal Agents Specific for P-glycoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/752,447 FILING DATE: 15-NOV-1996 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: 3'UTR
LOCATION: 4265..4669
SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: cDNA FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Mechetner, Eugene
                                                                                                                                                                                                                                                                                                                                   2454 GCTTGTGATCCACGGACACTCCTAC 2430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                   25;
                                                                                                                                                                                                                                                                                                                                                                           1 GCTTGTGATCCACGGACACTCCTAC 25
                                                                                                                                                                                                                                                                                                                                                                                                                                       h 100.0%; Score 25; DB 18; Length 4669; Similarity 100.0%; Pred. No. 0.027;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Chicago
STATE: Illinois
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 4669 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff Ltd
STREET: 300 South Wacker Drive, Seventh Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CHARACTERISTICS:
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425..4264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5'UTR
1..424
                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
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US-09-306-417-2/c
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                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 2, Application US/09306417 Patent No. US20020103144A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/306,417
CURRENT FILING DATE: 1999-05-06
EARLIER APPLICATION NUMBER: DE 198 22 115
EARLIER FILING DATE: 1998-05-08
NUMBER OF SEQ ID NOS: 17
                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Heinrich-Pette-Institut
TITLE OF INVENTION: Retroviral Gene Transfer Vectors
FILE REFERENCE: P50491
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: misc feature
LOCATION: (5775)..(8630)
OTHER INFORMATION: plasmid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: misc_feature LOCATION: (161)..(677)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LOCATION: (1)...(160)
OTHER INFORMATION: plasmid backbone (pUC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: (1)...(8630)
OTHER INFORMATION: retroviral expression vector SFbeta71m4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature
LOCATION: (1)..(8630)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
NAME/KEY: misc_feature
LOCATION: (5215)..(5774)
OTHER INFORMATION: 3'-LTR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LOCATION: (1220)...(5062)
OTHER INFORMATION: m4 mdr-1 cDNA
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OTHER INFORMATION: 5'-LTR
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LOCATION: (532)..(1219)
                                                          NAME/KEY: misc_feature
LOCATION: (1)..(8630)
OTHER INFORMATION: retroviral
                                                                                                                                      OTHER INFORMATION: Description of Artificial Sequence: proviral OTHER INFORMATION: plasmid DNA
                                                                                                                                                                                                                       TYPE: DNA
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AAD39004 standard; DNA; | | | | 100.0 | | | | | | | | | • | | | | • | ٠ | • | | • | • | • | 100.0 | • | 100.0 | |
| | l; DNA; | | 53099 | 8630 | 8630 | 6505 | 5544 | 4669 | 4669 | 4669 | 4669 | 4646 | 4646 | 4646 | 4646 | 4646 | 4646 | 4646 | 4646 | 4643 | 4643 | 4643 | 4643 | 4643 | 4643 | 4643 | 4533 | |
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| | | | Absystas Human mur | 1 Ketro | | | 4 Human | CDNA enco | | | | 7 | | Adk60994 Ovarian c | | | | | | μ | | Human | CDNA 6 | Abv74349 Human ABC | | CDNA enc | | |

Human; haematologic malignancy; multidrug resistance; MDR; SUMO-1; hypoxia inducible factor-1; small ubiquitin-like-modifier; HIF-1; lymphotid disorder; chronic lymphoproliferative disorder; lymphoma; myeloid disorder; lymphocytic leukaemia; thrombocythaemia; myeloma; angiogenic myeloid metaplasia; myeloid leukaemia; gene therapy; angiogenic wera; hypoxia responsive element; HRE; antisense; Human mdr1-HRE antisense oligonucleotide #7. 25-OCT-2001; 02-MAY-2002. Synthetic. Homo sapiens. Colgan 26-OCT-2000; WO200234291-A2 modified_base phosphorothioate (BGHM) BRIGHAM & WOMENS HOSPITAL INC. 2000US-0243542P 2001WO-US049856. Location/Qualifiers /*tag= a /mod_base= OTHER backbone; note= "Phosphorothioate backbone" 88.

Treating a subject (at risk of) having a hematologic malignancy or multidrug resistance, e.g. lymphoma or myeloma, by administering hypoxia

WPI; 2002-471427/50.

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RESULT 2
AAD39014/c
ID AAD390
XX AAD390
XX AAD390
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TX Human;
KW Hympho
KW Mypoxi
PT Misc_L
PT Misc_L
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; haematologic malignancy; multidrug resistance; MDR; SUMO-1; hypoxia inducible factor-1; small ubiquitin-like-modifier; HIF-1; lymphoid disorder; chronic lymphoproliferative disorder; lymphoma; myeloid disorder; lymphocytic leukaemia; thrombocythaemia; myeloma; angiogenic myeloid metaplasia; myeloid leukaemia; gene therapy; polycythaemia vera; hypoxia responsive element; HRE; ds.
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                                       Treating a subject (at
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                                                                                WPI; 2002-471427/50
                                                                                                                                                                                                           26-OCT-2000; 2000US-0243542P
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               risk of) having a hematologic malignancy e.g. lymphoma or myeloma, by administering
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multidrug resistance, e.g. lymphoma

hypoxia.

The present invention relates to a standardized polynucleotide system, which comprises at least one carrier nucleic acid, at least 3 oligonucleotides, as primers and target-gecific, fluorescently labeled probe and optionally at least one set of stabilized controls (standard RNA or DNA) of known concentration and instructions. The system comprise any of 20 sets of one control, two primers and one target-specific probe

comprises

Claim 1; Page 12; 38pp;

German.

standards,

primers

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ADK66076/c
ID ADK660
XX ADK660
XX O6-MAY
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XX Stuniden
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                                                                                                                                                                                                                                                                                         Standardized polynucleotide system, useful for quantitative, determination of nucleic acid, comprises stabilized standard
                                                                                                                                                                                                                                                                                                                                                                                                                                    Koehler T,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    functional genomics; sample analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ROBO-) ROBOSCREEN GES MOLEKULARE BIOTECHNOLOGIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standardized polynucleotide system; medical diagnosis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Rost
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llarity 100.0%;
Conservative (
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Pred. No. 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       pharmacogenomics; sample analysis
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RESULT 4
AAAO70907/c
ID AAQ70907 standard; DNA; 2726 BP
XX
AC AAQ70907;
XX
AC AAQ70907;
XX
DT 25-MAR-2003 (revised)
DT 27-MAR-1995 (first entry)
XX
Multidrug-resistance; MDR-1 gen
XX
Multidrug-resistance; MDR-1 gen
XX
Multidrug-resistance; MDR-1 gen
XX
VA Multidrug-resistance; MDR-1 gen
XX
CO 4-AUG-1994.
XX
PN W09417206-A1.

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Best Local S
Matches 25
                                                                                                        Query Match
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Matches 25
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                                                                                                                                                                                                                                                                      This gene is used as an example of a target DNA in a method for the amplification of nucleic acid sequences in a mixture using various DNA probes/primers. The method may be used to detect a mutation or allele ithe target, e.g. that that causes a genetic disease. (Updated on 25-MAI 2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Enzymatic amplification of target nucleic acid sequences in a mixt. detect a mutation or allele in the target, e.g. that causes a genetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 30; 50pp; English
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534
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25; Conser
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GCTTGTGATCCACGGACACTCCTAC 510
                                                    GCTTGTGATCCACGGACACTCCTAC 25
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llarity 100.0%;
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Pred. No.
                                                                                                                                                                                                                          C; 642
                                                                                                                                     Score 25; DB 2;
Pred. No. 0.081;
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RESULT 5 AAQ70916/c

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RESULT 6
AAT43326
ID AAT433
XX AAT433
XX O1-SEP
XX PCR; p
XX PCR; p
XX LCR; h
XX LCR; h
XX Synthe
XX WO9639
PD 12-DEC
XX X
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Best Local (
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                                                                                                                                                                                                                                                                                                                                            This gene is used as an example of a target DNA in a method for the amplification of nucleic acid sequences in a mixture using various DNA probes/primers. The method may be used to detect a mutation or allele the target, e.g. that that causes a genetic disease. (Opdated on 25-MA) 2003 to correct PN field.)
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16-DEC-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
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27-MAR-1995
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                                                                                                                                                                                           AAT43322 standard; DNA;
                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page.30; 69pp; English.
  05-JUN-1995;
                   04-JUN-1996;
                                                                               Synthetic.
                                                                                                                              Multidrug resistance gene-1
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                                      12-DEC-1996
                                                           WO9639537-A1
                                                                                                 primer; amplify;
human; multidrug
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l Similarity 100.0%;
25; Conservative 0
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93US-00168621.
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  95US-00461823
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resistance
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Pred. No.
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gene;
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                                                                                                   reaction; ligase chain reaction; MDR-1; ds.
                                                                                                                                                                                                                                                                                                                            728 T;
                                                                                                                                                                                                                                                                                              0.081;
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                                                                                                                                                                                                                                                                                      0
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RESULT 7
ABV78146/c
ID ABV781
XX ABV781
XX ABV781
XX Human
XX RNA ir
KW Viruci
XX Homo E
XX Homo
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Best Local S
Matches 25
                                 09-JAN-2001; 2001DE-01000586.
26-0CT-2001; 2001DE-01055280.
29-NOV-2001; 2001DE-01058411.
07-DEC-2001; 2001DE-01060151.
                                                                                                                                                                                                                                                                                                                            WO200255693-A2
                                                                                                                                                                                         09-JAN-2002; 2002WO-EP000152
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human mdr-1
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to be known
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                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
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protozoacide; gene; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Double stranded RNA; dsRNA; RNAi; RNA inhibition; cytostatic; viru protozoacide; gene expression; antisense; tumour; infection; Plasm virus; viroid; anti-GFP; human; HIV; human immunodeficiency virus; Hepatitis C virus; human papilloma virus; gene; ds.
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Inhibiting expression of target genes, useful by introducing into cells two double-stranded to the target.

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for treating tumors, that are complements

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Kreutzer

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Claim 13; Page 27-28; 100pp; German

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RESULT 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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This invention describes a novel method for inhibiting expression of a target gene by introducing into the cell that contains the target gene at least one oligoribonucleotide (dsRNAI) that has a double-stranded (ds) structure of not more than 49 consecutive nucleotides (nt), where at least a segment of one strand of the ds structure is complementary with the target gene and the cells are treatted with interferon before introduction of dsRNAI. The method is used to inhibit expression of target genes, particularly oncogenes, cytokine genes, id (not defined) protein genes, developmental or prion genes, or genes expressed in pathogenic organisms (particularly plasmodia) or in viruses or viroids (pathogenic in humans, animals or plants). Treating the cells with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oligoribonucleotide; interferon; oncogene; cytokine; Id; developmental; prion; inhibition; human; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Limmer S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      738 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
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Pred. No. 0.083;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hadwiger
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 3840;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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GCTTGTGATCCACGGACACTCCTAC

Query Match Best Local S Matches 25

Similarity

100.0%; ilarity 100.0%; Conservative 0

<u>,</u>

Score 25; DB 6; Pred. No. 0.083;); Mismatches

6

Length 3840; Indels

0;

Gaps

0

Sequence

3840

B₽;

1130 A; 738 C; 957 G;

1015 T; 0 U; 0 Other;

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RESULT 10
ABL91687/c
ID ABL916
XX ABL916
XX ABL916
XX ABL916
XX ABL916
XX Human
DE Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         문
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   interferon greatly increases the extent to which dsRNA can inhibit expression of the target genes, and the effect is even greater when dsRNA are modified to increase their stability. ABX09936-ABX10075 represent gene fragments used to illustrate the method of the invention
                                                   The invention relates to a method for inhibiting expression of a target gene (ABL91658-ABL91797) in a cell by introducing at least one oligoribonucleotide that has a double-stranded structure consisting of at most 49 sequential nucleotide pairs, with at least part of one strand complementary with the target gene and has at least one end a single-stranded segment of 1-4 nt. The method provides oligoribonucleotides for antisense inhibition of gene expression useful e.g. for treating tumours but the oligoribonucleotides may also be directed against genes present in pathogens (e.g. Plasmodium or viruses/viroids, pathogenic on humans, animals or plants) or against cytokine, Id, developmental or prion genes. The method provides more effective inhibition of gene expression than use of known oligonucleotides, probably because the unpaired overhang increases stability and thus intracellular concentration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; HIV; HCV; gene expression; oligoribonucleotide; tumour; pathogen; plasmodium; virus; viroid; cytokine; prion; antisense oligonucleotide; cytostatic; virucide; protozoacide; antibacterial; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 3840 BP; 1130 A; 738 C; 957 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human polynucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABL91687;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 13; Page 28-30; 104pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Inhibiting gene expression in cells, useful for e.g. treating tumors, introducing double-stranded complementary oligoRNA having unpaired
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-270454/32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kreutzer R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-JAN-2001; 2001DE-01000586
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (RIBO-) RIBOPHARMA AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Limmer S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA; 3840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1015
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   T; 0 U; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
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RESULT 12
AAZ49332/c
ID AAZ493
XX
AC AAZ493
AC AAZ493
XX
DT 14-MAI
XX
DE Human
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ABO78185/c
ID 818781
XX ABQ781
XX ABQ781
XX YX Human;
XX Human;
XX Homo s
XX Homo s
XX Homo s
XX Homo s
XX IFT CDS
FT CDS
FT Variat
FT AUG
FT AUG
XX JP2002

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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to presumption of the side effect of at least one immunosuppressant selected from tacrolimus and cyclosporin in which if the 2677th base in the position of MDR1 gene in the encoding region of the cDNA sequence is guanine, adenine or thymine, is investigated. The method is used for the presumption of the side effect of an immunosuppressant. The present sequence is that of the MDR1 encoding cDNA of the invention
                                 Human wild-type multidrug resistance-1 (MDR-1) cDNA.
                                                                                                                                     AAZ49332;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; immunosuppressant; tacrolimus;
single nucleotide polymorphism; gene;
                                                                                   14-MAR-2000
                                                                                                                                                                                    AAZ49332 standard; cDNA; 3860
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 3843 BP; 1129 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 6-11; 14pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       primer set.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Presuming the side effect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31-JAN-2001; 2001JP-00024723
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-JAN-2001; 2001JP-00024723
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                                                                                                                                                                                                                                                                                                              2030
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DB; ABB83950.
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
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                                                                                                                                                                                                                                                                                                                                              GCTTGTGATCCACGGACACTCCTAC 25
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                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           replace (2677, A/T)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /standard_name= "Single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /product= "MDR1"
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                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 25; DB 6; 100.0%; Pred. No. 0.083;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  740 C; 956 G;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ID NO 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1018 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 σ,
                                                                                                                                                                                                                                                                                                                                                                                                                  <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 3843;
                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
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resistance protein MDR-1 MDR-1 is a transmembrane efflux pump, cresponsible for the export of drugs from cells, particularly cancer cells. Wild-type MDR-1 shows increased resistance to etoposide and decreased resistance to etoposide and convertion relates to transducing haematopoietic stem cells with nucleic acid encoding an MDR protein and culturing the modified cells. The modified haematopoietic stem cells are useful in bone marrow transplantation (to reconstitute haematopoietic systems in patients who therapy of genetic defects in cells derived from haematopoietic stem cells are useful and in ex vivo gene therapy of genetic defects in cells derived from haematopoietic stem cells, e.g., thalassaemia, Gauchier's disease, sickle cell anaemia or cells, e.g., thalassaemia, Gauchier's disease, sickle cell anaemia or leukaemia. The modified cells can also be used to identify factors involved in regulating proliferation and differentiation in haematopoietic stem cells that express MDR-1 haematopoietic stem cells that express MDR-1 will be provided and the contrafted cells.
will be protected against chemotherapeutic agents, so can be engrafted while the patient is undergoing chemotherapy. Expansion of (rare) haematopoietic stem cells provides sufficient cells to permit standard biochemical analysis. Overexpression of MDR-1 allows cytokine-driven expansion of haematopoietic stem cells by at least 10-fold compared with a maximum of 4-fold in known procedures
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This sequence represents cDNA encoding human wild-type multidrug resistance protein MDR-1. MDR-1 is a transmembrane efflux pump,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 10; Page 68-70; 113pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ex vivo expansion of hematopoietic stem cells transduced with a sequence encoding human multidrug resistance-1, used for bone marrow
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sorrentino
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                02-DEC-1999
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/note= "cDNA sequence of G185V
in AAZ49333"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /*tag= a
/product= "Human wild-type MDR-1 protein"
replace(553. .555, GTT)
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2030 GCTTGTGATCCACGGACACTCCTAC

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GCTTGTGATCCACGGACACTCCTAC 25

Query Match Best Local S Matches

Similarity

100.0%;

Score 25; Pred. No. C; 957 G;

DB 3; 1022 H

> Length 3860; U; 0 Other;

0

Conservative

0;

Mismatches

0

Indels

0

Gaps

0

Sequence 3860

BP; 1135 A; 746

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AAZ49333/c

AAZ4933XX

AC AAZ493

XX AAZ493

XX Human

XX Synthe

OS Synthe

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OS Synthe

OS Synthe

XX Synthe

OS OS This self

OS This self

OS This self

OS This self

OS C This
                                                                                                                                                                           CC resistance protein MDR-1, where the the Gly residue at postion 185 of the cefflux pump, responsible for the export of drugs from cells, particularly CC cancer cells. The wild-type MDR-1 shows increased resistance to etoposide and decreased resistance to vinca alkaloids compared with the G185V CC mutant. The invention relates to transducing haematopoietic stem cells CC with nucleic acid encoding an MDR protein and culturing the modified with the G185V CC cells. The modified haematopoietic stem cells are useful in bone marrow CC transplantation (to reconstitute haematopoietic systems in patients who CC have undergone chemotherapy or radiation therapy) and in ex vivo gene CC cells, e.g., thalassaemia, Gauchier's disease, sickle cell anaemia or CC leukaemia. The modified cells can also be used to identify factors CC involved in regulating proliferation and differentiation in haematopoietic stem cells that express MDR-1 CC will be protected against chemotherapeutic agents, so can be engrafted CC while the patient is undergoing chemotherapy. Expansion of (rare) thaematopoietic stem cells that express MDR-1 CC manatopoietic stem cells to permit standard CC biochemical analysis. Overexpression of MDR-1 allows cytokine-driven cCC expansion of haematopoietic stem cells by at least 10-fold compared with cc a maximum of 4-fold in known procedures
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Query Match
Best Local Similarity
Matches 25; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Multidrug resistance; MDR-1; P-glycoprotein; transmembrane efflux pump; haematopoietic stem cell; transduction; bone marrow transplantation; chemotherapy; radiation therapy; cancer; gene therapy; gene replacement; genetic defect; thalassaemia; Gauchler's disease; sickle cell anaemia; leukaemia; ex vivo expansion; cytokine; mutant; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ex vivo expansion of hematopoietic stem cells transduced with a sequence encoding human multidrug resistance-1, used for bone marrow
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-DEC-1999.
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                                                                                                                               Sequence 3860
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 79-82; 113pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       transplantation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (SJUD-) ST JUDE CHILDREN'S RES HOSPITAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9961589-A2
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   100.0%;
llarity 100.0%;
Conservative 0;
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1. .3843
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/product= "Human G185V mutant MDR-1 protein"
                                                                                                                               1135 A; 746 C; 956 G; 1023 T; 0 U; 0 Other;
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   Score 25; DB 3; Length 3860;
Pred. No. 0.083;
); Mismatches 0; Indels
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      Gaps
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ARBSULT 14
ARBS435/C
IID ARBS433
XX ARBS433
XX ARBS433
XX ARBS433
XX ARBS433
XX ARBS433
XX ARBS43
DT 26-MAR
XX Stem C
KW hepato
KW cardia
XX Key cardia
XX Key C
FH CDS
FT CDS
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밁
                                                  Ś
                                                                                                           Matches
                                                                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                            disorder, myocardial infarction, Parkinson's disease, degenerative disorders of the brain, and for tissue regeneration or replacement. Haematopoietic cells can be used in bone marrow transplants (e.g., for treatment of leukemia) and for ex vivo gene therapy for treating blood diseases such as sickle cell anemia and thalassemia. The stem cells can also be used as cell targets in gene therapy protocols. The present sequence represents a sequence related to the BCRP for which no relevant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hepatotropic; n cardiant; gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Stem cell; ATP transport protein; ATP-binding cassette; antiparkinsonian; hepatotropic; neurodegenerative; cytostatic; antianemic; muscular; BCRP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human BCRP DNA related seq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-MAR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention provides a method of identifying and/or isolating a ste-
cell that involves detecting the expression of an ATP transport protection on the provided and the containing a conserved ATP-binding cassette (BCRP) by a cell in a san comprising stem cells. The isolated stem cells may be used in the treatment of diseases such as muscular dystrophy, degenerative liver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Identifying a stem cell, for treating e.g., muscular dystrophy, myocardial infarction, Parkinson's disease, or neurodegenerative disorders, comprises detecting the expression of an ATP transport protein (BCRP) by a cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
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29-MAY-2001;
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                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 53-55; 87pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sorrentino B,
                                                                                                                                                                                                                                                                      information has been provided in the specification
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                                                                                                           25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABB07266.
                                                                                                                                       Similarity
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  GCTTGTGATCCACGGACACTCCTAC
                                                        GCTTGTGATCCACGGACACTCCTAC 25
                                                                                                                                                                                                                      3860
                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                     The invention provides a method of identifying and/or isolating a stem cell that involves detecting the expression of an ATP transport protein containing a conserved ATP-binding cassette (BCRP) by a cell in a sample comprising stem cells. The isolated stem cells may be used in the treatment of diseases such as muscular dystrophy, degenerative liver disorder, myocardial infarction, Parkinson's disease, degenerative disorders of the brain, and for tissue regeneration or replacement. Haematopoietic cells can be used in bone marrow transplants (e.g., for treatment of leukemia) and for ex vivo gene therapy for treating blood diseases such as sickle cell anemia and thalassemia. The stem cells can also be used as cell targets in gene therapy protocols. The present seguence represents a seguence related to the BCRP for which no relevant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Identifying a stem cell, for treating e.g., muscular dystrophy, myocardial infarction, Parkinson's disease, or neurodegenerative disorders, comprises detecting the expression of an ATP transport protein
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P-PSDB; ABB07267.
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29-MAY-2001; 2001US-00866866.
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ALIGNMENTS

JOURNAL COMMENT ACCESSION VERSION KEYWORDS SOURCE REFERENCE AUTHORS LOCUS DEFINITION RESULT 1 BP243819/c 밁 ORIGIN FEATURES TITLE Query Match 100.0%; Score 25; DB 5; Best Local Similarity 100.0%; Pred. No. 0.56; Matches 25; Conservative 0; Mismatches 0; ORGANISM source Homo sapiens (human) Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo. El (bases 1 to 568) Suzuki,Y., Yamashita,R., Shirota,M., Sakakibara,Y., Chiba,J., Mizushima-Sugano,J., Nakai,K. and Sugano,S. Sequence comparison of human and mouse genes reveals a homologous block structure in the promoter regions Genome Res. 14 (9), 1711-1718 (2004) Contact: Yutaka Suzuki Department of Virology Institute of Medical Science, University of Tokyo 4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan Email: ysuzuki@ims.u-tokyo.ac.jp. BP243819 Sugano cDNA library, liver HepG2 Homo sapiens cDNA clone HEP21483, mRNA sequence. BP243819 BP243819 BP243819 GI:52116729 EST. /organism="Homo sapiens" /mol type="mRNA" /db Xref="reaxon:9606" /clone="HEP21483" /tissue type="liver" /cell line="HepG2" /clone lib="Sugano cDNA library, liver HepG2" /note="hepatoma" ocation/Qualifiers Length 568; Indels <u>,</u> Gaps 0;

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Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp.
Location/Qualifiers
                                                                                                                                      Clark,A.G., Glanowski.S., Nielson,R., Thomas,P., Kejariwal,A.,
Todd,M.A., Tanenbaum,D.M., Civello,D.R., Lu,F., Murphy,B.,
Ferriera,S., Wang,G., Zheng,X.H., White,T.J., Sninsky,J.J.,
Adams,M.D. and Cargill,M.
                                                                                                                                                                                                                                                                                                                 Clark, A.G., Glanowski, S., Nielson, R., Thomas, P., Kejariwal, A., Todd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F., Murphy, B., Ferriera, S., Wang, G., Zheng, X.H., White, T.J., Sninsky, J.J., Adams, M.D. and Cargill, M.
Inferring nonneutral evolution from human-chimp-mouse orthologous
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1 (bases 1 to 579)

Suzuki,Y., Yamashita,R., Shirota,M., Sakakibara,Y., Chiba,J., Mizushima-Sugano,J., Nakai,K. and Sugano,S.

Sequence comparison of human and mouse genes reveals a homologous block structure in the promoter regions
Genome Res. 14 (9), 1711-1718 (2004)

Contact: Yutaka Suzuki
                               Rockville, MD 20850, USA
This sequence was made by sequencing genomic exons and ordering them based on alignment.
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BP242809.1 GI:52115719
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                                                                                                                             2030
BX645450 497 bp
DKFZp781C1343_r1 781 (synonym: h
DKFZp781C1343_5', mRNA sequence.
                                                                                                                                                                                              25;
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Clark, A.G., Glanowski, S., Nielson, R., Thomas, P., Kejariwal, A., Todd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F., Murphy, B., Ferriera, S., Wang, G., Zheng, X.H., White, T.J., Sninsky, J.J., Adams, M.D. and Cargill, M.
Direct Submission
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Clark, A.G., Glanowski, S., Nielson, R., Thomas, P., Kejariwal, A., Todd, M.A., Tanenbaum, D.M., Civello, D.R., Lu, F., Murphy, B., Ferriera, S., Wang, G., Zheng, X.H., White, T.J., Sninsky, J.J., Adams, M.D. and Cargill, M.
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AY408955
AY408955.1 GI:39764923
GSS.
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25; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (16-NOV-2003) Celera Genomics, Rockville, MD 20850, USA
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                                                                                                                                                                                                                                                                                                                               /mol_type="genomic_DN
/db_xref="taxon:9598"
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                      /gene="ABCB1"
                                                                                                                                                                                                                                                                                                                                                                     organism="Pan troglodytes"
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/locus_tag="HCM3396"
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                                                                                                                                                                                                           100.0%; Score 25; DB 9
100.0%; Pred. No. 0.63;
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                   497 bp mRNA
lonym: hlcc4) Homo
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VIRTUAL TRANSCRIPT, partial sequence,
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                   CDNA clone
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This clone (DKFzp781C1343) is available at the RZPD in Berlin.
Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ingolstaedter Landstr.1, D-85764 Neuherberg, Germany This is the 5' sequence of the clone insert Clone from S. Wiemann, Molecular Genome Analysis, German Cancer Research Center (DKFZ); Email s.wiemann@dkfz- heidelberg.de; sequenced by DKFZ (German Cancer Research Center, heidelberg/Germany) within the cDNA sequencing consortium of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Wellenreuther, R.,
Unpublished (2003)
Contact: MIPS
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1 (bases 1 to 497)

Poustka,A., Albert,R., Moosmayer,P., Schupp,I., Wellenreuther,R., Mewes,H.W., Weil,B., Amid,C., Osanger,A., Fobo,G., Han,M. and
                                                                                                                                                                                                                                                                                                                                                         534 bp mRNA linear EST 01-JAN-EST01407 Atlantic salmon Lambda Zap Express ovary cDNA library Salmo salar cDNA clone oyr 004 d08 5', mXNA sequence.

BM414475
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BX645450.1 GI:34479783
Contact: Caplice NC Department of Microbiology National University of Ireland Galway (NUIG) National University of Ireland Galway, Galway Tel: 00353 91 524411 2254 Fax: 00353 91 525700
                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Actinopterygii; Neopterygii; Teleosteei; Euteleostei; Protacanthopterygii; Salmoniformes; Salmonidae; Salmo. 1 (bases 1 to 534)
Caplice, N.C., Martin, S.A., Davey, G.C. and Powell, R. An expressed sequence tag-based list of genes expressed in the testis and overy of Atlantic salmon (Salmo salar)
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                                                                                                                  Unpublished (2002)
                                                                                                                                                                                                                                                                                                Salmo salar (Atlantic salmon)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone_lib="781 (synonym: hlcc4)"
/note="Vector: pSport1_Sfi; Site_1: SfiIA; Site_2: SfiIB;
cDNA-collection"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /db xref="taxon:9606"
/clone="DKFZp781C1343"
/dev_stage="adult"
/lab_host="DH10B"
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|mol_type="mRNA"
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100.0%; Pred. No.
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                                          Galway, Ireland
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356 CTTGTGATCCAGGTACACTCCTAC 333
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Plate: ovary rare plate 4 row: d column: 08

Seq primer: M13 reverse primer = caggaaacagctatgacc
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22; Conserv
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Construction and primary characterization of normalized cDNA
libraries in rainbow trout, Oncorhynchus mykiss
Unpublished (2003)
On Apr 7, 2003 this sequence version replaced gi:29598411.
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BX317766 tcay Oncorhynchus mykiss

BX317766 tcay Oncorhynchus mykiss
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                                                                                                                                                                   Email: Yann.Guiguen@beaulieu.rennes.inra.fr
Sequence cleaned of vector, adaptator and re
at sigenasupport@jouy.inra.fr to obtain the
                                                                                                                                                                                                                                                 Campus de beaulieu, RENNES
Tel: 02.23.48.50.09
Fax: 02.23.48.50.20
                                                                                                                                                                                                                                                                                                                       On Apr 7, 2003 this
Contact: Guiguen Y
INRA - SCRIBE
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1 (bases 1 to 569)
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EcoR1; Site 2: Xho1; An Atlantic salmon ovary cDNA library
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/db_xref="taxon:8030"
/clone="oyr_004_d08"
1. .569
/organism="Oncorhynchus mykiss"
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/mol_type="mRNA"
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/dev_stage="adult"
/clone_lib="Atlantic salmon Lambda Zap Express ovary cDNA
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                                                                Location/Qualifiers
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Pred. No. 56;
0; Mismatches
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RESULT 8
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Construction and primary characterization of normalized cDNA
libraries in rainbow trout, Oncorhynchus mykiss
Unpublished (2003)
On Apr 7, 2003 this sequence version replaced gi:29600852.
Contact: Guiguen Y
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BX320207
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: Yann.Guiguen@beaulieu.rennes.inra.fr
Sequence cleaned of vector, adaptator and re
at sigenasupport@jouy.inra.fr to obtain the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Campus de beaulieu, RENNES cedex, 35042, Tel: 02.23.48.50.09
Fax: 02.23.48.50.20
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1 (bases 1 to 574)
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/tissue_type="adipose tissue, blood, brain,
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differentiating gonads, gills, interrenal, intestine,
kidney, liver, muscle, ovary, pituitary, testis"
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/dev_stage="from embryos to adults"
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/clone_lib="tcay"
/clone_lib="tcay"
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normalized + 1 subtraction (tcay); Clone distribution :
AGENAE Resource centre. Francois PIUMI,
Francois.Piumi@jouy.inra.fr, INRA, CEA Radiobiologie et
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone lib="tcay"
/note="Vector: pT7T3D-pac; Rainbow trout multi-tissues -
/note="Vector: pT7T3D-pac; Rainbow trout multi-tissues -
normalized + 1 subtraction (tcay); Clone distribution :
AGENAE Resource centre. Francois PIUMI,
AGENAE Resource feature. Francois PIUMI,
Francois Plumi@jouy.inra.fr, INRA, CEA Radiobiologie et
Francois Plumi@jouy.inra.fr, INRA, CEA Radiobiologie et
Etude du genome (IREG), Domaine de Vilvert, 78352,
Etude du genome (IREG), Domaine de Vilvert, 78352,
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/tissue_type="adipose tissue, blood, brain,
differentiating gonads, gills, interrenal, intestine,
kidney, liver, muscle, ovary, pituitary, testis"
/dev_stage="from embryos to adults"
/lab_host="phi03"
                                                                                                                                                                                                                                                                                                                                                /mol_type="mRNA"
/db_xref="taxon:8022"
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                                                                                                                                                                                                                                                                                                                                                                                                        organism="Oncorhynchus mykiss"
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/db_xref="taxon:8022"
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REFERENCE AUTHORS TITLE

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FEATURES

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RESULT 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: bkoop@wic.ca
Genome Sciences Centre, BC Cancer Agency
Genome Sciences Centre, BC Cancer Agency
CDNA preparation, sequencing and bioinformatics:
CDNA preparation, sequencing and bioinformatics:
Y Butterfield, R Kirkpatrick, J Asano, N Girn, R Guin, D Le
S Lee, T Olson, P Pandoh, A Prahbu, D Smailus, L Spence, J
S Taylor, G Yang, J Schein, S Jones and M Marra.
Location/Qualifiers
ssalmge003036 gut Salmo salar
CA044923
CA044923.1 GI:24349786
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University of Victoria
PO Box 3020 STN CSC, Victoria
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A. survey of Salmo salar transcripts from high complexity cDNA
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Salmo salar
                                                         CA044923
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Contact: Koop BF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tel: 250 472 4067 Fax: 250 472 4075
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                                                                                                                                                                                                                                                                                                                                   /clone lib="mixed_tissue"
/clone lib="mixed_tissue"
/note="Vector: pCMVsport6; Library Creator: Research
/note="Vector: pCMVsport6; Library Creator: Carlo
Genetics; Atlantic salmon tissue contributors: Carlo
Biagi, Mitch Uh and Robert Devlin (DFO, Vancouver, B.C.)
Simon Jones (PBS, Nanaimo, B.C.), Seaspring Hatchery
(Crofton, B.C.), Rachel Roper (University of Victoria)"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        organism="Salmo salar"
/mol_type="mRNA"
/strain="McConnell"
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Email: bkoop@uvic.ca
Centre for Biomedical Research, University of Victoria
cDNA preparation and sequencing:
cDNA preparation and sequencing:
Roberto Alberto, Marianne Beetz-Sargent, Maura Busby,
Roberto Hunt. Linda McKinnel, BF Koop.
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Salmo salar (Atlantic salmon)
Salmo salar
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1 (bases 1 to 592)
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Fax: 250 472 4075
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University of Victoria
PO Box 3020 STN CSC, Victoria BC,
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                                                                                                                                                                        Oncorhynchus mykiss (rainbow trout)
Oncorhynchus mykiss
Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei;
Protacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus.
1 (bases 1 to 595)
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                                                                                                                            Govoroun, M., Guiguen, Y. and Le Gac, F. Construction and primary characterization of normalized cDNA libraries in rainbow trout, Oncorhynchus mykiss
                                                                                                                                                                                                                                                                                                          mRNA sequence.
BX910867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      bioinformatics:
                                                                                           Unpublished (2003)
On Jan 22, 2004 this sequence version replaced gi:41127666
                                                                                                                                                                                                                                                                                           BX910867.2 GI:43415782
               Tel: 02.23.48.50.09
Fax: 02.23.48.50.20
                                               Campus de beaulieu,
                                                              Contact: Guiguen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
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Yann.Guiguen@beaulieu.rennes.inra.fr
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/strain="McConnell"
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Pred. No. 56;
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Campus de beaulieu, I
Tel: 02.23.48.50.09
Fax: 02.23.48.50.20
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Seq primer: M13R.
                                                                                                                                                                                                                                                                  Email: Yann.Guiguen@beaulieu.rennes.inra.fr
Sequence cleaned of vector, adaptator and repetitions. Contact us
at sigenasupport@jouy.inra.fr to obtain the chromatogram of this
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Govoroun, M., Guiguen, Y. and Le Gac, F.
Construction and primary characterization of normalized cDNA
libraries in rainbow trout, Oncorhynchus mykiss
Unpublished (2003)
On Dec 17, 2003 this sequence version replaced gi:39992052.
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/tissue_type="multi-tissues"
/dev_stage="from embryos to adults"
/lab_host="DH108"
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/mol_type="mRNA"
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/mol_type="mRNA"
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cDNA preparation, sequencing and bioinformatics:
Y Butterfield, R Kirkpatrick, J Asano, N Girn, R Guin, D
S Lee, T Olson, P Pandoh, A Prabbu, D Smailus, L Spence,
S Taylor, G Yang, J Schein, S Jones and M Marra.
Location/Qualifiers
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ssalrgb531310 mixed_tissue Salmo
CA052009
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Genome Sciences Centre,
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University of Victoria
PO Box 3020 STN CSC, Victoria BC,
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A survey of Salmo salar transcripts from high complexity cDNA
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Proteacanthopterygii; Salmoniformes; Salmonidae; Salmo.
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/clone lib="mixed_tissue"
/note="Vector: pcWVsport6; Library Creator: Research
Genetics; Atlantic salmon tissue contributors: Carlo
Biagi, Mitch Uh and Robert Devlin (DFO, Vancouver, B.C.),
Simon Jones (PBS, Nanaimo, B.C.), Seaspring Hatchery
(Crofton, B.C.), Rachel Roper (University of Victoria)"
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Francois.Piumi@jouy.inra.fr, INRA, CEA Radiobiologie et
Etude du genome (LREG), Domaine de Vilvert, 78352,
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/mol_type="mRNA"
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Salmo salar
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Contact: Koop BF
                      Unpublished (2002)
                                                               http://web.uvic.ca/cbr/grasp
A survey of Salmo salar trans
                                                                                                      GRASP Consortium, Davidson, W.S., Koop, B.F. and
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Sequence cleaned of vector, adaptator and repetitions. Contact
at sigenasupport@jouy.inra.fr to obtain the chromatogram of th
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Fax: 02.23.48.50.20
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Construction and primary characterization of normalized cDNA
Libraries in rainbow trout, Oncorhynchus mykiss
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Oncorhynchus mykiss
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                                             libraries
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/mol type="mRNA"
/db_xref="texon:8022"
/clone="tcbi0021c.n.22"
/tissue_type="testis"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Resource centre. Francois PIUMI, Francois.Piumi@jouy.inra.fr, INRA, CEA Radiobiologie et Etude du genome (LREG), Domaine de Vilvert, 78352, Jouy-en-Josas cedex, FRANCE, +33 (0) 1.34.65.28.02, +33 (0) 1.34.65.22.73"
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91.7%;
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                                                                      2 CTTGTGATCCACGGACACTCCTAC 25
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University of Victoria PC V8W 3N5, Canada
PO Box 3020 STN CSC, Victoria BC, V8W 3N5, Canada
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Genome Sciences Centre, BC Cancer Agency cDNA preparation,
Genuencing and bioinformatics: Y Butterfield, R Kirkpatrick, J
sequencing and bioinformatics: Y Butterfield, R Kirkpatrick, J
sequencing and bioinformatics: Y Butterfield, R Kirkpatrick, J
smallus, L Spence, J Stott, S Taylor, G Yang, J Schein, S Jones and
M Marra.
POLYA-Yes.
                                                                                                                                           83.2%; Score 20.8; DB 6; llarity 91.7%; Pred. No. 56; Conservative 0; Mismatches 2;
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